

749

AGTCCGGGGG ACCTTTTAG TCGGTAGATT GAGATTGCAA ACAAATCTGC ATCTACATTG	4860
AAAGCTTAAT TTCTAATAAT TGAAAAAATC GAATGAAAAA TTTCTTACCT TCATTCACAG	4920
AGCTCGATTT CAGAGCTCTT TTTGCTAGCT TATTCTACT TTTCTGAATT TCGAAAAAGA	4980
AATGTAAGCG TTTGATAGAT TTACAAAAAG ATTGTATAAT AGGGATAAGA ATAGAAAAGG	5040
AGAAGTCTCA TGGCAGTTAA AGATTTATG ACCCGCAAGG TAGTTTATAT TAGTCCAGAT	5100
ATAACAGTAT CTCATGCAGC AGATTTGATG AGAGAGCAAG GTTGCACCG TCTGCCTGTT	5160
ATCGAAAATG ATCAATTAGT TGGTTGGTG ACTGAGGGAA CCATTGCACA AGCAAGTCCA	5220
TCTAAAGCAA CAAGTCTTTC TATCTATGAG ATGAATTATC TTCTGAATAA GACAAAAGTA	5280
AAAGATGTCA TGATTCGCGA TGTTGTCACT GTCTCAGGCT ATGCTAGTCT AGAAGATGCA	5340
ACTTATCTGA TGTTGAAAAA TAAGATTAGT ATTCTCCCTG TCGTAGATAA CCATCAAGTA	5400
TACGGAGTTA TTACTGACCG TGACGTTTC CAAGCCTTC TTGAAATTGC AGGTTATGGC	5460
GAAGAAGGGA TTCGTGTACG CTTTGTACA GAAGATGAAG TTGGTGTCT TGGAAAAATT	5520
GTTTCTTGA TTGTTAGAAGA AAATTGAAT ATCTCCCATA CAGTCAATAT TCCGCGTAAG	5580
GATGGTAAGG TGATTATCGA AGTGCCTAAC GATGGATCAA TTGATTTACC AGCCTTGAAA	5640
GAAAATTTG AAGCAAATGG TATTCAAGTG GAAGAAATCG CTCGCACCTTC AGCAAAAGTC	5700
TTGTAAGAAG GGAAGCCAA AGGCTCTTT TTTCATGAAA AGGGGATTAG AGCAAAAGAT	5760
GGAAAGAAAT GATAAAATAT GCTATAATGA AATAATGTAA AAAAGGAGTA TTTATGGACA	5820
TTTCAGTAAT TCGTCAGAAA ATTGACGCAA ATCGTAAAA ATTAGCTTCT TTCAGGGGT	5880
CTCTTTGACC TCGAAGGGCT AGAGGAAGAG ATTGCCATCT TGGAAAACAA GATGACAGAA	5940
CCTGATTTTT GGAACGATAA TATTGGGCC CAAAAAACGT CGCAAGAATT AAATGAATT	6000
AAAAACACTT ACAATACCTT CCATAAGATG GAAGAGTTGC AGGATGAAGT CGAAATTAA	6060
TTGGATTTTT TGGCTGAAGA CGAGTCAGTG CATGATGAAC TGGTAGCGCA GTAGCCGAA	6120
CTTGATAAGA TAATGACCAAG CTACGAGATG ACTCTACTCT TGTCAGAAC TTATGACCA	6180
AACAATGCCA TCTTGGAAAT CCATCCAGGT TCTGGTGGTA CTGAGGCGCA GGACTGGGT	6240
GATATGTTGC TTCGTATGTA TACTCGTTAT GGTAAATGCTA AAGGCTTAA AGTGGAAAGTG	6300
TTGGATTACC AAGCAGGTGA TGAGGCTGGT ATTAAGTCGG TAACTTTATC ATTTGAAGGG	6360
CCTAATGCCT ATGGTCTCCT CAAGTCAGAA ATGGGTGTTG ACCGCTTAGT GCGAATCTCA	6420
CCATTTGACT CTGCCAACG TCGCCATACC TCTTTCACAT CTGTAGAAGT GATGCCAGAA	6480
TTGGATGATA CTATTGAAGT GGAAATCCGT GAAGATGATA TCAAGATGGA TACCTCCGT	6540

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TCAGGTGGTG	CCGGTGGACA	AAACGTCAAT	AAGGTTCAA	CAGGTGTACG	TTTAACCCAC	6600	
ATTCCAAC	TG	GAATTGTTGT	CCAATCAACA	GTAGATCGTA	CCCAGTATGG	AAATAGAGAT	6660
CGTGC	CATGA	AGATGTTGCA	GGCTAAGCTC	TATCAAATGG	AGCAAGATAA	GAAGGCTGCG	6720
GAGGTAGATT	CTCTCAAAGG	TGAGAAAAAG	GAGATCACTT	GGGGAAAGCA	AATCCGTTCT	6780	
TATGTCTTCA	CGCCTTATAC	TATGGTAAA	GATCACCGAA	CTAGCTTGA	GGTTGCTCAG	6840	
GTAGATAAGG	TTATGGATGG	GGACCTAGAT	GGTTTATCG	ATGCTTATCT	CAAGTGGCGA	6900	
ATTAGCTAAG	ATAGAAAGGA	ACTCACATGT	CAATTATTGA	AATGAGAGAT	GTCGTTAAA	6960	
AATACGACAA	CGGAACA	ACT GCTCTACGCG	GTGTTCGGT	TAGCGTTCAA	CCGGGGAAAT	7020	
TTGCTTACAT	CGTAGGACCT	TCAGGAGCAG	GGAAGTCAAC	TTTTATTCTG	TCTCTGTATC	7080	
GTGAAGTAAA	AATCGATAAA	GGAAGCCTAT	CAGTTGCTGG	TTTTAATCTG	GTAAAGATCA	7140	
AAAAGAAAGA	TGTCCCCTT	CTACGTCGA	GTGTTGGGT	TGTCTCCAG	GATTATAAA	7200	
TGTTACCAAA	GAAA	ACTGTC	TATGAAAATA	TTGCTTACGC	TATGGAAGTA	ATCGGGAAA	7260
ATCGCCGTA	A	TATCAAAAGA	CGAGTGATGG	AAGTTTGG	CTTGGTTGG	TTGAAGCATA	7320
AGGTTCGTTC	TTTCCC	AAAT GAACTCTCAG	GTGGGGAGCA	ACAGCGGATT	GCGATTGCGC	7380	
GTGCAATTGT	AAATAATCCC	AAAGTATTGA	TAGCTGATGA	GCCAACAGGA	AATCTGGATC	7440	
CGGATAATT	TC	ATGGAAATT	ATGAATCTCT	TGGAACGGAT	TAACyTACAA	GGAACAACTA	7500
TTTTGATGGC	GA	CTCAT	AAAT AGCCAGATTG	TAAATACCTT	GCGCCACC	GTGATTGCCA	7560
TTGAAAATGG	CCG	TGTCGTT	CGTGACGAAT	CAAAGGAGA	GTATGGATAC	GATGATTAGT	7620
AGATTTTTTC	GCC	ATTATT	TGAAGCCTTA	AAAAGTTGA	AACGAAATGG	TTGGATGACA	7680
GTAGCTGCTG	TCAG	TTCAGT	CATGATTACT	TTGACCTTGG	TGGCAATATT	TGCATCTGTT	7740
ATTTTCATA	CAG	CGAAACT	AGCTACAGAT	ATTGAAAATA	ATGTCG	TGT AGTTAT	7800
ATCCGAAAGG	ATG	TGGAAGA	TAATAGTCAG	ACAATTGAA	AAGAAGGTCA	AACTGTTACA	7860
AATAATGACT	ACC	ACAAGGT	ATATGATTCT	TTGAAGAAC	TGTCTACGGT	AAAAGTGTT	7920
ACCTTTCAA	GTAAAGAAGA	ACAATATGAA	AAATTAAACCG	AGATAATGG	AGATAACTGG	7980	
AAAATCTTG	AAGGAGATGC	CAATCCTCTC	TATGATGCCT	ATATTG	TAGA	GGCAAACACT	8040
CCAAATGATG	TAAA	AAACTAT	AGCCGAAGAT	GCTAAAAAAA	TTGAAGGTGT	CTCTGAGGTT	8100
CAAGATGGCG	GTG	CCAATAC	AGAAAGACTC	TTCAAGTTAG	CTTCATTTAT	CCGTGTTGG	8160
GGACTAGGGA	TTG	CTGCTTT	TTAATT	ATCGCAGTT	TCTTGATTTC	AAATACCATT	8220
CGTATTACCA	TTAT	TTCCCG	CAGTCGCGAA	ATTCAAATCA	TGCGCTTGGT	CGGAGCTAAA	8280
AACAGTTATA	TCC	GTGGACC	GTTCTGTTA	GAAGGAGCCT	TTATCGGTTT	ATTGGGAGCT	8340

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ATCGCACCAT	CTGTTTGTT	CTTTATTGTT	TATCAAATTG	TTTACCAATC	TGTCAACAAA	8400
TCGTTGGTAG	GGCAAAATCT	ATCCATGATT	AGTCCAGATT	TATTTAGTCC	GTTGATGATT	8460
GCCCTACTAT	TTGTGATTGG	GGTTTCATT	GGTCATTGG	GATCAGGAAT	ATCCATGCGC	8520
CGATTCTTGA	AGATTTAGGT	AAAATAGCTG	CTTTTATGAG	GAGATTGTAA	AATCTCCTTT	8580
TTTGCTACAA	GAGTTTTGA	AAAGAGATGC	GCAGAAGAAA	AGAGCTTCCA	AAGAAGTCCC	8640
CCAGAGAAGA	CTTC					8654

(2) INFORMATION FOR SEQ ID NO: 99:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 19718 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 99:

TGTCGCGTCA	AAATCATTAC	TATGGCTATG	TATAGCCCTT	ACTATGACTT	GGCTAACAC	60
GTTCGCTTTC	AAATTCTAG	GCTCAGGCTG	AAACAGTCTC	CCAGGCTGTT	CACTCCGAA	120
TGCTAAAATC	GTTCTTGATC	GCTTTCACAT	TGTACAACAT	CTTAGCCGTG	CTATGAGTCG	180
TGTGCATGTC	CAAATCATGA	ATCAGTTCA	TCGAAATCC	CATGAATAACA	AGGCTATCAA	240
GCGCTACTGG	AAACTCATTC	AACAGGATAG	CCGTAAACTG	AGTGATAAGC	GATTTATCG	300
CCCTACTTTT	CGCATGCACT	TAACAAATAA	AGAAATTCTT	GACAAGATTT	TAAGCTATTC	360
AGAAGACTTG	AAACACCACT	ATCAGATCTA	TCAACTCTTA	CTTTTTCACT	TTCAGAACAA	420
AGACCCTGAG	AAATTTTCG	GACTCATTGA	GGACAATCTG	AAGCAGGTTTC	ATCCTCTTT	480
TCAGACTGTC	TTTAAAACCT	TTCTCAAAGA	TAAAGAAAAG	ATTATCAACG	CCCTCAACT	540
ACACTATTCT	AATGCCAAC	TGGAAGCGAC	CAATAATCTC	ATCAAACCTTA	TCAAGCGCAA	600
TGCCTTGGT	TTTCGAAACT	TTGAAAACCT	CAAAAAACGG	ATTTTTATCG	CTTTGAACAT	660
CAAAAAAGAA	AGGACGAAAT	TTGTCCTTTC	TCGAGCTTAG	CTGACTTCAA	CCCACATACAG	720
TTGACAAAGA	GCCTAATTTC	CATAAAATT	GACATGGAAA	TTATAAAACC	ATTACTAGTT	780
TAGTCCTTTT	TGATAACGTG	CCAATTCCGC	TTGGTTCGCC	CAAACATAGT	GACCTGGACG	840
GATTTCCTACC	ATAGATGGCT	TATCAGTCTC	ATAGTCGTGT	TGACTTGGAT	CGTAAACCTT	900
CAAGACCTTC	TTACGTTCCA	AGATTGGATC	TGGGATTGGT	ACCGCTGAAA	GCAAGGCTTG	960
AGTATATGGG	TGAATTGGAT	TGTTAACAA	TTCTCTGTT	TCTGCAACCT	CTACAATAAC	1020

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ACCCCTTGAA	ATAACTGCGA	TACGATCTGA	AATAAAGCGA	ACAACCGACA	AGTCATGGGC	1080
GATGAAGAGA	TAGGTCAGGC	CGAGCTCTT	TTGGAATT	TTGAGCAAGT	TCAAGACTTG	1140
GGCACGTACA	GAAACGTCCA	AGGCTGAAAT	TGGCTCATCT	GCAATAACAA	AGTCTGGTTG	1200
CATGACCAAG	GCACGGCAA	TACCAGATCG	TTGACGTTGA	CCGCCTGAGA	ATTCACTGAGG	1260
GTAACGAGTC	AAGTGCCTAG	CAAGAAGACC	TACTTCACGG	ATAATATTT	GAACCTTCTC	1320
TTTACGTTCT	TCTTCATCCT	AAAATAACG	GTGATTGTAA	AGACCTTCAG	AAATAATATA	1380
ATCAACAGTC	GCACGTTCAT	TCAAACTTGC	GGCAGGGTCT	TGGAAAATCA	TCTGGATTG	1440
ACGAATCAAT	TCCGCACTT	GTTCACGCGA	TTTCTTACCA	TTAATCTTT	GACCATCAAA	1500
AATGATATCT	CCATTACTTG	TATCATTAG	ACCGATGATA	GCACGACCAA	TAGTTGTTTT	1560
CCCACTACCG	GACTCACCTA	CAAGCGAGAA	AGTTTCTCCC	TTGTTGATAA	AGAAGTTAGC	1620
ATTTTTAAC	GCGACAAACT	TCTTACTTCC	TTCACCGAAG	GAAATTCTA	AATCTTGAT	1680
TTCTACTAAT	TTTCAGACA	TTTCCTTCCT	CCTAGTCAGC	CAGATGGC	AATCCCATTT	1740
TTTCACGGAT	CTTATCATGG	AGATTTGCAA	TCACAGCTGG	TTTTTCTACT	TTCGGAGCAT	1800
CCTCATGAAG	AAGCCAAGTT	TTAGCCCAAT	GTGTCTCTGA	TACTGAGAAT	TGAGGAGCTT	1860
TTTGTTCGAA	GTCATCTGC	ATPGCGTAGT	CAGAACGCAA	GGCAAAAGCA	TCCCCTTCA	1920
GGTCAGTATA	AAGTGACGGA	GGTGTCCCTG	GGATTGAGTA	AAGATCCCCT	TTATCATCAG	1980
CAAGCTGAGG	CAAGCTAGAC	AAGAGACTCC	ATGTATATGG	ATGGCGAGGG	TCATAGAAGA	2040
CTTCCTCAAC	CGTTCCATAC	TCAACGATTT	CTCCTGCATA	CATAACCGCT	ACCTTATCCG	2100
CAATACTTGC	CACCAACACCA	AGGTGCTGGG	TAATAAAGAT	TGTTGTAAA	TGATACTCGT	2160
TTTGTAAAGA	TTTTAGCAA	TCAATAATCT	GAGCTTGAAT	AGTTACATCC	AAGGCAGTTG	2220
TTGGCTCATC	ACAGATCAAG	ACATCAGGTC	GGCAGGCAAG	GGCAATAGCA	ATAACGATAC	2280
GTTGACGCAT	TCCTCCAGAA	TATTGGAATG	GGTATTCTATT	AAAACGTCTA	TCTGCGTCTG	2340
GAATGCCAAC	CTTATTCTATG	TAGTCAATGG	CCAATTCTTT	CGCTTCTTTA	GCTGTTTTTC	2400
CTTGGTGT	TACAATAACT	TCTGTAATCT	GACTACCAAT	TGTTTTAATG	GGGTCCAAC	2460
TAGTCATTGG	GTCCTGGAAG	ATAGTCGCAA	TCTTAGCACC	ACGAATTGT	TCCCAATCCT	2520
TGTGAGAAGA	TAAAGCTGTC	AAGTCCTGAC	CACGGTAGTC	AATACTACCT	TGGGCAATAC	2580
GACCATTTC	TTCGAGCATA	CCTGTGAAGG	TCTTGTCAA	AACAGATT	CTGATCCTG	2640
ACTCACCTAC	CAAGGCTAAT	ACTTCTCCTT	CGACTAGTTC	AAGGGAAACG	CCGCGAATGG	2700
CTGTCAATAC	TTTGTACGAA	ACGTCAAATT	CCACGACAAT	ATCGCGAGCA	GTCAAAATT	2760
CATTTTTTC	TTTTGTCATT	TCTACTCCTA	TCTATGTGTA	CGTGGATCAC	TAGCATCCGC	2820

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TAAGTTTGAA	CCAAC TACGA	AAAGGGACAA	GGATACCAAG	ACAAGGGTTG	TCAATGGAAT	2880	
CCAGAACAAAG	TAAGCATTGG	TTGTTACGTT	TTGTGAATAA	TCCGAAATCA	AACGACCCAA	2940	
ACTTGGCACT	GTAATCGGT	ATCCAAGACC	GAAGAAAAGAC	AAGAAGGCCTT	CGTATGAGAT	3000	
AAAGCTTGGGA	AGCATTGAG	TCATGGTTGT	CACAATAACA	GATACCAATT	GAGGCATGAT	3060	
ATTTTTGGCA	ACAATCTCA	AGGTTGGTGT	TCCCCAAAGTA	CGTGACGCCA	AGTTGTATT	3120	
CAAGTCACGA	TAGCGCAAGA	TTTGCACACG	GATCATGAAG	GCAATACCAA	TCCATGTTGT	3180	
TACGCTCATG	GC AAAAATCA	GATTCCAGAA	TCCAGCTCCG	ATTGAGTAAG	TCAAGACAAT	3240	
AAACATCAAA	AGAGGTGGGA	TGTTTGAGAT	GACGTTGTAA	ACTTCCATCA	TGACACGGTC	3300	
AACTGATT	T GAAATACCCC	AAATACCACC	GACAAAAAAC	CCGATAACCA	AGTTAAC	3360	
TGTCGCAATC	ACAGAAAATGA	GGATGGAGTT	ACGAGCTCCG	AACCAGACAC	CGTCAAAGAG	3420	
CGATTTACCG	TTACTGTCAG	TACCGAACCA	ATGCTCCGCA	TTTGGCTTGA	TATAACGAAC	3480	
ACTAAAGTCG	TTTACCTTGC	TGACATCATT	GAAATCAAAC	TTAGAAAACA	TTGGGTAGAT	3540	
GAAACTTATC	AAAATGATGG	CTACCAAGAT	TCCCAACATG	ACTACAGTTG	ATTTTTCTT	3600	
CATAAAATTGT	TTAAACACTG	ATTTCAGTA	AGAATATGCT	GGCGCATCAA	TAGTTTCAGA	3660	
GGCAAATCG	TCACGTTTA	CAAAC TGAAA	TTTTCTTTA	TCGATTGTAG	ACATTATTG	3720	
CCTCCTTCT	CAGTCATT	AATACGTGGG	TCAATAATAG	TCATCCAAAT	ATCTCCAAA	3780	
AGACGTGAGA	AGATAGAAAT	ACATGTAAAG	ATGAAGACAA	GACCAACGAC	CATAGAGTTA	3840	
TTAGATGCTT	TTACAGAGTC	AATCAACATT	TTACCCATAC	CTGGGAAGGC	GAAGACTGTT	3900	
TCAGTAAGGG	TTGCACCAACC	GATAACCCCA	ATAATGGCAG	CAGGAATTCC	TGAAACCAGC	3960	
GGAACCATGG	CATTTTAAA	GATGTGTTG	TTTGAAATT	CTTTTCAGA	CAAACCTTT	4020	
GCACGAGCGA	AAAGAACAAA	GTCTTGAGAT	TGCAAGTCAA	TCATGTAACG	ACGAATCAA	4080	
ATGGCTGTAC	CAGGAGCACC	CAACAAACCA	AGGATGACTG	CTGGTAAAAC	GTAAGAACGC	4140	
CAATCTCCAG	CTCCCAAGAT	AGGGAATGAA	TCTGGAAGGG	CAATAGATGA	TCCAATCAA	4200	
CGAACGATGT	AAACCAAGGC	AATCGTTGGA	AGAGCAAGCA	AGAAGGTCAA	AGCCCCTGTT	4260	
GAGAGGCTAT	CAATCCAAGT	GTTCTTGAAA	CGAGCCATGG	CTGAACCAAG	TGGCACGGCA	4320	
AGAGCATAGG	CAAGAACCAA	ACCAATCAA	CCAGTAATAG	CAGAGCTGAC	AATCATAGAT	4380	
GGATATTGGT	AATTACTTTC	AGTCGCTGTA	TAAGGATCAT	CTTTCCCATA	GCTAGCTACT	4440	
TCACGAGAGT	CAGCCTGACT	AGGTGACTTG	TAGGTTCTTG	AGTAAATATT	TACAGAAC	4500	
GT	TTTCTTAC	CTGTTGGAA	CTGAACCTGG	GCAGTTTG	TTTGTCCCTG	ACCTTGAGTA	4560

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ATAACCTGAA	GAACCTGGTGT	ATTAGCATAG	GTTGGGTAAG	AGTCACCTAA	ATTCAAGTTC	4620
ACAAAGTTT	GATGAACAAA	TGGGAACGTGA	CTGTTAAAGT	ACAAGAGATA	TTTATGTTA	4680
GTTCCCTGAAC	CGACCAATGA	CCATCCGATA	GCTGGATCAT	TTTCAAAACG	AAGGTAGCGT	4740
TTCAAGTCTG	GATTTTCAGG	GTCTTGATT	TTATTTGTAT	GGTCAATGTC	AATCAAGTTA	4800
GCATAGAACGT	AAAAAACACG	TTCAAAAATT	GGAATTCAC	GAGTAGCATA	GAATTGACCA	4860
CTTTCAGTAA	ATTCTCCAA	AGTCCAACCA	TGACCTAATT	GATTGATGTA	CTTTTCATAAA	4920
ATAGCTTAT	TGGTCGCATT	TGCTTCTACT	GTTACAGAAG	AATCCATGCT	ACTTGCCTTT	4980
TCTTGCAACT	CTTTAGTATC	GTAATACTCA	ATGTAGCCCA	TACGCTCAA	CACAGTATTT	5040
TCATAGTTAT	CACGTTTATC	AGCCGTTGTC	GCAATTTCAT	TATAGTTAGG	ATCCTGCTTG	5100
AAAATCAATT	TTCGAGGAAC	CAAGGTATAG	ATAATCGTGT	AGGTCAAAGT	CGTTACTAAG	5160
AAAATCGAAA	CCAATGACCG	CAAAACACGC	ATAAAAATAT	ATTTTTTCAT	ATTATTTCC	5220
TTAAAAATCC	CAAAAGAACCC	TTCTCCTCAT	GGAGAGAAAG	TTCTATTAGA	AATTATTTAC	5280
TTCACATGAC	TTGCCAATTC	TTTTTGAGCT	TTCTCATTG	ATTCAGCTTT	TTCTTTCAAC	5340
CATTTTCAC	GAGCTTTTC	ATACTCTTCC	TTAGTCACCA	CTTTATCTTG	TGATTTCAAA	5400
TATTTGAAGT	AAACATCTGA	CCCCTTAGAG	CCTGTTGCG	CAGAAGCTCC	AGTAAATGGA	5460
ACAATTCTGTG	AAAGCACTGG	TGCTGCACCA	GAAGAAGCCA	TAGCAGGAAT	AAAGAGTGAA	5520
CTATCTGTCA	ACCATGCTTG	AGCCGCTGCA	TATTTTCAT	AACGGACATT	CAAGTCGCTT	5580
GTCTCTCTGG	CAGCTTCATC	AACTAATTAA	TCGTATTCTT	TCAAACCAAC	TTGAACTACT	5640
GAAGGGCTAT	TTGGATTATC	AAATCCTAAA	TATGTTTTG	TAGTTTCACT	GCTAGTTGTT	5700
TTTAAAATAT	CCAGGTAAGT	AGATGGGTCT	TGATAGTCTG	GCCCCCATGA	AACTCCTCCT	5760
GATACATCCC	AATCCTCAGA	TGAAGCATTG	GCAGCATAGT	AAGTAATATT	AAGGAATTCA	5820
TCACTTGTCA	TTTGTGAAAT	ATCAACAAACG	ACATTTCAA	CACCAAGAAC	TGTTTCTACA	5880
GATTGTTAA	AGGACTGAAT	ACGAGATATG	TAGTTTTTG	ATGCTTGTC	TACTGGAACG	5940
TCCAGATGAA	TAGGAAACTG	AACGCCGTCT	GCTTCTAAAG	CTTTCTTAGC	TTTCGCAAAC	6000
TCTGCCTTGG	CCTTGTCA	ATTGAATAAA	CCATCCTGCC	CATCAGCTAA	ATTCACACCT	6060
TTCCACTCAT	CACCATAAGC	AGGAAGTTGA	GCAGCGACTA	AATCACCAAA	GGTCTCTCA	6120
CCAGCTGAAA	CAAAGCTCTGG	TTTACAAAT	AAATTACGAA	CTGCTAAAGC	TGCTCCATCT	6180
TTACCATTGA	TTTGAGCTGA	GTAAGCTGAG	CGATCAAGAG	CAAATTCAA	GGCTTGACGG	6240
AAATCTTGT	TAAGCAATGC	CTTCTTAGTA	GCTACTTTCT	CTGAATCTGT	AGTTTAGAA	6300
GTATAGTTGT	AACTTTGGCG	ATCAATATTC	ACACCCAGAC	CAGCAATCCC	AGAGCCTGAT	6360

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TGTGTGTAAT AGATATTGTC CTTGTATTCT TCTGCAACCT TAGAATAGTT GGAGCTGGTA	6420
GGGTAAAGAC GGGCATAACT ATAAGCTCCA CTAGTGAAGT TACGCTCTAG CGACTCCTGA	6480
TCTGATCCAT CATAGTAAGC TAGATTGATA GTATCTAGGT GGACATTTC TTTATCCCAA	6540
TATTGCTCAT TTTTACAAA CTCTACAGAA GATTTGCAG TCAACCCTT CAACAAGAAT	6600
GGACCATTAT AAAGCAAGGA TGCGGATCT GTTGGTTAG CAAAATCGCT TCCTTTGAT	6660
GTTTCGAATT CCTCATTCAAG AGGCCAGAAA ATAGAATAGG TCAACTTAGA GTTCCAGAAC	6720
GGTTCAAGT GTATTGTAAC GTATAATCAT CAACCGCCTT GACACCAACT	6780
GTTGAAAAT CTGTTGAAGT TCCTGATAGA TAATCTGCCA AGCCTTAAAC CGAATTTC	6840
GCTAAATACA TAGCTCTGA TTTTTATCT GCTGCGTGT TTAAACCGTT CACGAAATCT	6900
TTAGCCGTCA CCTCTGCATA TTCTCTCCA TCAGAGGTA ACCATTAAAC CCCTTACGA	6960
ATCTTATAAG TGTAGGTCAA ACCATCCTTA GAGACTTCCC AATCCTCTGC AACTGCAGGA	7020
GCAAGATTAC CGTAATTATC GTTAGTGAAT AAACCATCAA TCCCATTGTA AGTCACTACT	7080
GTTGTACTAT TTTTACTTGA AATCAGGTAG TCCAAGGTTT CTGGGTCTGC TGTATAAAC	7140
TAGCCATAAG CTTTAGGGC TGATGAATCA GATGATTTG AAGAACTGCA TGCTGCAAGT	7200
ACACCTGCTG CTAATAAAAC AAGACCTGCT GTAGCAAATA CACGATTTT TTTCATTTC	7260
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ATCAAACAAA TTTTCAGAAT ATTTAGGCTT GTGGCACAA ATTTTTCTATT TTTTTGAAT	7380
ATATGATTCA AATTGTCGTT CGAAGTGTCA AAGACTACAG TGAAAATAGG AAATTGACG	7440
CAGAAACTTT GGAGTTAGG AAGACATACA GTAAAATGAA ATACGGACGG AACAAATGTGA	7500
TTTTGGAATT CAAATTAAAT TATAACAATA TTGTAGAAGT ATCATTCTAG TATTCAAGAT	7560
TCAGTTTACT ATGTCTTTTC ACACCAACCT TATCCGAAT TCAATTACTT TTGTGATTAA	7620
CATATATAGA TTAAGACTAT CTTTATACT TTAAAATTC TCGCTACCTT ATCCACTATA	7680
TGCTCCTCGC TATCACGTTT CTATTCAAG CCTACGATTT CACTATTGCT TTCTCTGACA	7740
ATTCTTATTTC CCTGCGTCAG ACTTAAAACG ATCTATCCCC AGACCATTAAATCCGCTAC	7800
CTCACGATAG TCAGGCTTGG GGAGCGCTAT TGTATTCACTT GGTAGTGGAG CCCTACAGAG	7860
GACTTACACC TCAGATGCAC GACATGCCA TCGTATAAAA AATCTCCTAC CCAAGGTAGA	7920
AGATTTCAAA CTTATAAAAC TTAATCCGTC ATGTCCGATA CCAACATTGCG ATGCTCCAAT	7980
GGAATACTGC ACATAACTAG CAAGAAAATA AAGCCTGACT GAATCCAGAA GAGAGCCAAG	8040
TCAAAATTC CGTGCACAGC AACCACTGTA AGGAAAGATA GATAAAGGCC GATAATCGGA	8100

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CGTTTCCCCG ACTCCTGACT CATATCCATC ATCAAGCGAA CAGGAGCAAC AGAAGACAAA	8160
ACTAATAAAA TAGTCCCCAC AATTCCGTAA CTCAGAATCG TATCAATATA AAGACTGTGG	8220
GCATGTTCAT GATAAGGAGC ATGTATCCGA GGATAAGAGT TCATATAGGT CAATGCCCT	8280
TCACCCCCAA AAGGATTTCG CTTAACAAAG GCCATCCCAG CATCCCAGAT AGAAATGCGT	8340
TCTTCCATAG AAGAGTCTAA AGTACCCATT CGAACTCCCA AATCACTAGA AAAGAGGAAA	8400
CTCAAACCAA TCGCGAAGAC CCCAATACTA AGCCAAAAGG CCTTCCAGTT TTTAATAGTC	8460
GTAAAGAGAT AGATAATTGC TCCAGCGATA ATAGCAGGAA AGGCAGTTCG ATTTTGAGTA	8520
AAGTTCAAAAC CAAAGAGATT AACAAAGCCT GCAATCACAC AGAATACTTT CAACCAATT	8580
AACTTGGTCG TTGTAACAG ATAGAAAGCA ATCATAATAC AGAAACACA AATAATTCCA	8640
TAATAATTAG GATTAAAGAA GGTCACTTCT GCCCGGTTCT GATGCCACAC CTGCATATTG	8700
GGTGAAAGAA AAGCATAGTT AAATTCTTC ACAATTGGA AATGTTCTAA ACTGGCAAAA	8760
GCAGCTGACA AGACACTACC AAACAAGACA AACTGCAAAA TCAATCGAAA GAATTATGG	8820
GATAAAATCG ACTGATAGT CAAAAAGAAA ATAGTAAATA GAAACATTCC TACTGAAGCC	8880
ACAAGACCCA TCCAATTTCG TGCAAGAATG GATATAACAG TACTATAGCT AAGAAAAGA	8940
AGCAGCATCG GATGCTCCC CATTTCCTGA AGAATACTTT TCATGTCTCC TGAAAAATC	9000
AAACTGATAA TATATAAACAA GAGTACAAC TACAAAAAGAT AAAAGGGTAA AAAGATACTC	9060
AGGATAATTTC CCAATAAAAT CAGCTCTTA CTAGACAAACC CCTTCAGCTT TTCAATAAG	9120
CCTATTGATT TCAAAATGAA TCCTTCTCT CCAAATCAGC TGATTCAGAT AATAGTAAGC	9180
TATCCTATAT TGTACCACTT TTTAGCAAT TTGAAAACAA AGGAAACGTT TTCCAAAATA	9240
AAAACCTAT TTTATCCACC ATATCAAGGC TTCAAAATGA TACTTCAACT CCATTCTCAA	9300
TTACCCGATA AGTCTGATTT TGCAAATCAA TTTCTACTAC TGCTGTTACG GACTTATCTT	9360
TATTTTGACG TTTGATTACA ATGCTGTGAG CTGTTGGTGT CTCTATCTCA GTAGTCCCTT	9420
CTAGATCAAA GGCTTCTGAA CGGTTACGGA AAGAAAATAG ATTGAGAAGG GCCTTCACAA	9480
CAGGTCGTTG CACTTCTTT GCTATTTCT CGTTGCTATA GTAATGACGA TTAATATTC	9540
GACCTTCTTT AGTTTCTCT AATAATTCA AGTCATTCTT GCCTGCTAAT AGACCCACAT	9600
AGTAAATCTG AGGAATAACCT GGGGCAAAAG CTTGAATTAG ACGAGCGAGA AAATACTTGA	9660
CATCATCATC TCCAAGCGCT GAATAGTAGG TTGAATTGAT TTGGTAGATA TCTAAGTTGT	9720
TATACTCGGC ACTAGAGTAC TTACGTTGA CATTGGCTCC AACCTTATAG AGTCATTTG	9780
AAGCATAGTC AATCTCCTCA TCGGTCAAGGA TATCCTTGAC ATCTACTACT CCAATCCCAT	9840
CATGGGTATC TAGCGTCGTA AATTGCTTCAT TCGGGCTCAT CTTTAACCAC TTAGCCAAAC	9900

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GCTCTGTTCT	GGAACGTAA	AGAGTATAAA	GTGTCACCAT	TGGAAGAGCA	AAATCATAAA	9960
CATAGTAATC	ATGGTCTGCT	ATTTTAAACT	GAATCGAATA	GTGTTCATGA	ATCTCAGGTA	10020
AAAGCTCTGT	CCCATACTCA	GCAGCGATAT	CTCGAACCTT	GTCCAATAAA	TCCCAAATAT	10080
CTGGTTCAC	AAAGAAATCA	TTAGTATCCA	ATTCTTCAC	TGCATAAGCA	AAGGCATCTA	10140
GACGAATCAA	ATCACACCCA	TTACTTGCCA	AGTGCTGAAT	GGTCTTACGG	ATAAATTCCA	10200
TAGTTACTTC	TTTGGTCACA	TCAAGATCAA	TCTGCTCCTC	ACCAAAGGTA	TTCCACAAAT	10260
GTTCCACTGA	ACCATCTTCA	AACACAATCT	CTTGCTTTGG	TGCACGATCC	TTACGCTTGT	10320
AAATTAAATC	TACATCAGAC	TGTGTCGGAC	GGTTTCTGG	CCAAAACCTA	TCCCAGTTA	10380
AAAAGAGAGC	TTTAAATTCA	CTGGCTTCAT	GTTTTCTTG	ATAGTCCTTA	TAATACTTGG	10440
ATTGACGAGA	AATATGATTA	ATCATAAAAT	CAAACATAAG	ATAATATTTC	TCACCTAAC	10500
GCTTCACATC	CTCCCAATCA	CCAAAAGCTG	AGTCCACTTC	GTCGTAGTCA	ACTGGCGCAA	10560
ATCCACGATC	AACTGTTGAT	GGGAAAAATG	GTAAAAGGTG	AACTCCTCCA	ATAGCATCTC	10620
CAAAATGCTC	TTCCAAATTA	TCATATAAGT	CTTAAAGATT	ATTTCAAGG	CTATCAGAAT	10680
AGGTAATCAA	CATGGTTTA	TTTGAAATTG	GCATCATTAC	TCTCCTTTT	CTAATTGAAG	10740
CCAAGTCTCA	TATGATCTGG	CTTCATAAAT	AAAATTCTT	TTAAATCTCT	ATTTATCATC	10800
AAACTCGTAC	TAATATAGAC	TGTGATAAAC	AAAGTACTAC	TTTCTTGTIT	TCTGCATAGA	10860
ATTATCAACA	AGCTAAACTC	TTCCCTCTGTG	TCAAAGACTA	TAGATTCCAT	GAGCTCTTCT	10920
TATACTCTTC	GAAAATCTCT	TCAAACCACG	TCAGCTTCAC	CTTGCCGTAG	GTATGGTTAC	10980
TGACTTCGTC	AGTTTCATCC	ACAACCTCAA	AACAGTGTTT	TGAGCAACCT	GCGGCTAGCT	11040
TCCTAGTTG	CTCTTTGATT	TTCATTGAGT	ATTACTTCAC	TGCCCGTTG	CTCATTCCCTG	11100
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AAGGCAGAGT	CCACATTTCG	GAATCCCGGT	TCAAGACAAG	GAGTGGCAAC	ATGAAGTCAT	11280
TCCAGAACCA	AAGGGCATTG	ATGATCATGG	TTGTCGCATG	CATCGGTTTC	ATCATTGGGA	11340
AGATGATGCC	GAAATAGGTT	GTAAATTGAT	TAGCCCCATC	GATCTCTGCT	GCTTCATCCA	11400
GACTTTCTGG	AATCGAGATT	TTGATATAGC	CAACATAGAG	AAAGAGGGTC	TGTGGAATCG	11460
CATAGGTCAA	GTAGAGCAAG	ATCAAACCAA	AGGTATTAGC	CAAACCGAGT	TTACTCATCA	11520
TAACCGTAAT	CGGAATCATG	ATGACTTGGG	AAGGTACGAA	GATTCCGAGG	ATTAAGAGGG	11580
TATACATGAT	GGTAAAGGCT	TTTCTTTAC	TCATATTGCG	AGCGATGGAG	TAGGCTGCCA	11640

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GAAAGCCAAA GAAATTATCT ACAATATCCT TAGTGGGTTT GAAGGAACTA AAGAGGGTAG	11820
CAAGGAGCGG CACTAAAATC AGAACCGATC CTAGAATCAA TAGAATGTAT TTGCCAATCA	11880
GGGCTTTCTC TTCATCTTGT TTCATCATGC TTCTCCTCTT AAATTCAAA TTTCTTAGAT	11940
ACTCTCAATT GGATGATCGA AACACTACA ATTAAGAAGA ACAAGATTAC GGCAATGGCA	12000
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GTTGTGGCAT TGTGGGACC ACCACCGGTC ATGGCAAAGA CTTGGTCAAAG GGCAGTCAGC	12120
CCACCTTTA GGGCTAGGAT AAAGACCATA GAGACACTTG GTAGCAAGTA AGGCAATTCA	12180
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GTTGGAATAG ATTGCAAACC AGCTAGGAAG ATGATGATGG GCATAGCCAC CCCTTGCCAA	12300
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AAAAATTCAA TATGAAGGGC ATTTCCAATC GCTGGAAGAC CGTAGTTGAA GACTTGCTTG	12420
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AAGGTTTGGC CTTTGATTT AGAATTCAAG ACACCGCAGA TGAAGATCCC GAGTGCAATC	12540
TCACCAACCA CCATGGCAAT CGCAATGATT GCGGTAAAGC CAATCGCATT CATGAATTTC	12600
GGATCCATGA AGAGGAGCTT AAAGTTGTTT AAGCCAACAA ATTTGTAGTT ATAAGTCAAT	12660
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GCTTGTAACA AGAGGGGAT GACCACAAAA GCCCATGCC AATATTTTG TAATACTTTT	12780
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GTCTGCTTCA CTGGTCCAGT ATTGTTGCAA CCAGACCAAG TGACGATCCG TAAAGGCATA	12960
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TGGAGATCCG TCCACATCGT AGTATTTTG CATGACTTCT GGACGGGTCA TATATTCCAC	13080
AAAGGCATTG GCTTCTTTG GATGTTGGT GGTGGCTGAG ATAGACCATG CCAAGTCTCC	13140
CGCACCAACG GTTAAGCTTT GTCCCTTTTC TTTTCCTGGA ATCATGAAGG TCCCAATCTT	13200
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TTGCTTAGAT CCATTGATGC GAAGGATGTC CATGACCTTG ATATCATCTT TCATAATCGG	13380
ATCCGACAAT TTAATGGCAT TTGGTTGAGA ATAACGAAGG TATTGATTG CTTCTTTCC	13440

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TCCACCTGTT	GCTGTCGCAA	AGGCTAATTG	ATTGTAACCA	TTGAGTGTCC	AAGCATCTGC	13500
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ACCTGCTTT	GCCCATTCTT	GCAGTTCGAT	GGACTGTGGG	TAAATATTGA	CCACATCAGG	13800
CACATCTCCT	GCGAGAACGC	GTGTCTTCAA	TACTTCACCA	GCATTTGGTA	CATTGACGAC	13860
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GGTCATTTCT	TTTTCTGGT	TGAAATACTC	GATGGTCACT	GTGCCATCCG	CAGATTACC	13980
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CACCCCCAAA	GTTAGACAGA	ATAAAATCTAA	CTTTTGGGGT	CAGTACATAT	CATAGTTTC	14160
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CGTAGGTTAC	AATCGTTGA	TTTCCGTAAT	TAAATTGTAC	AGCTGCTTCA	TTGGATAACAG	14400
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GTCCCATCTG	ATGATTGGT	ACTGCTGACA	CATGAGCCCC	CATAGAAATG	GTTGGATAGA	14640
GATAGGATGA	ACCGTATTGA	ATTGGTAAAC	GTGCAATGCC	ATCAGTATTA	TCACTAGCCC	14700
AGACTTGTGG	GAAATAGCGC	ATCATACCAA	GATCATTCG	TCCACCACCA	CCAGAGCAGG	14760
ACTCAAAGAG	AATATGGCTG	TGCTTCTCTG	TCAGATAAGA	AACGAGTTCA	TAAAGCCCCA	14820
GCATGTACTG	ATGAGATTGC	ATCTGTGTCT	CTAGATAAGT	TAATCCATT	CCTAGCTTAG	14880
TGATATTGCG	GTTCATATCC	CATTTAATGT	AATCAATATC	ATGATAAAAT	AGGAGTTGAT	14940
CTAAGACACT	TTTCAAGTAT	TCTACTACCT	GAGGATTGGC	AAGATTAAGT	ACTAATTGAT	15000
TCCGAGAATA	AGTATGCTA	TAGCCAGGAA	CCTGAATAGC	CCAGTCAGGA	TGTTGACGAT	15060
ACAAATCACT	ATCTACAGAA	ATCATTTCGG	GTTCTAACCA	AAGTCCAAAC	TGCAAACCTC	15120
TTTCATGGAT	AGCTGAAATC	AGACTTCTA	GACTTCCACC	CAGTTTTCC	TCATTAACAA	15180

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CCCAATCACC TAAAGCACGA TTATCATCAA AACGATTGCC AAACCAACCA TCATCTAATA	15240
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GAAAGTCAAA GTAAGTAGCT TCCCAGTTAT TGATTAGAAT TGGACGTTCT TTTTAGAAA	15360
ATTCACTTAG CATAATGTGC TTCAGTACAA ATTCTGACT TTCACTGACTA ATACCAGTTA	15420
ATCCCTGATC TGAATGAGTC ACTAAAGCTA CCGGTGTTTC AAAGTATTCC TCAGGAGCTA	15480
ACTTCCAAGA AAAGTTTCT GGATTAATGC CAATAGCCAC CCGAAGTCA TTCAATTGAT	15540
TTTTTGAAC AAAAGCTTCA AAGTTGCCAC TATACATTAG TTGAATAGCA AACACATTCC	15600
CAGCATCCTC TGTGACTCCT TGTCGCATA GTAGAAGAGC TGGTGTGTTGA GCATGACCAG	15660
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TCTTTTCACG AGCATAAGCA CCCTGCAGAG TTACTATTTG GTAATCTGCA GCTGGAAAAT	15780
CAGCCATAAA AGAAAATCT TTATGGATGA CAACTTCCTG ATTACTATTA TTATCTAATT	15840
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AGTCTGTTAC TTCAGTTACA CTATGCTGAA CCTGTATGGT TGGTTCCCTA AAATCTCCTA	16080
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CCTCATCACT TTATTGATTA TATTTTATCA CCTGAAATCG CTTTCCAAAA TAGAAAATG	16380
TCTCAAGAAT ATGGTAAAAT GTTAGGTTAGG AGGTAGCACA TGTTAGTTTT TTCAGAATAC	16440
CAGACTGGAA CAATCGACCT TGCCCTAAGC TTTTATGGAT ATGAGGAATG CACACCTAAT	16500
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TTAGGAATCA CTGGAGGGAA AGCCCTGAT TATTTGCTC TTTCCAAAT TTCTGATCAA	16740
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CAAACCTCATG AACTGATGTT TCATCTGGGA ACTATTGCTC CCAATCAGAA AAAAAAGAAT	16920
ATTTCATCAA CCCACCAACT CTATCTTGAA TGCAACAGAT TAATTGATAG CCACTATCCT	16980

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CAATCACTTA	CAATTCAAGA	TTTAGCAAAA	GAACTATCCG	TTCACAGAAG	CTACTTATCA	17040
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CCAAGTCATA	CAAGAAAAGA	ATACTCTCAA	TACCAACTAG	TAAGAAAGGC	AACATTATGA	17280
AATCCTACCA	AGCTGTCTAC	CAAATCCTAT	CTAAAGAAC	CGACTATATC	AGCGGAGAAA	17340
AAATCGCAGA	AAAACATATCC	CTAAGCCGAA	CAGCAATTG	GAAAGCCATC	AAGCGACTAG	17400
AACAAGAAGG	CATTGAAATT	GATAGTATCA	AAAATAGAGG	ATATAAACTG	ATGAATGGTG	17460
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AAACAAAATC	AACACAACTA	GATGCAAAAG	AAGCAATTGA	TTTAGGCCAT	GAAGCAAATA	17580
CCCTCTATCT	AGCTTCCTAT	CAAACAGCAG	GCCGAGGCCG	TTTCAACGT	TCCTTCTACT	17640
CACCACAAGG	TGGTATTCTAT	ATGACACTCC	ATCTTAAACC	AAATCTCCCC	TATGACAAAT	17700
TACCATCCTA	CACACTACTT	GTAGCTGGAG	CTGTCTACAA	AGCCATTAAG	AACCTAACTT	17760
TAATAGATGT	CGACATAAAA	TGGGTCATG	ATATCTATCT	AAACAATCAT	AAAATTGGAG	17820
GAATCCTTAC	TGAAGCAATG	ACCTCTGTAG	AAACTGGCTT	AGTCACAGAT	ATCATTATTG	17880
GAGTAGGTAT	CAATTCACT	ATTAAAGACT	TCCCTCAGGA	ATTAAAAGAA	AAAGCTGCCA	17940
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TAGGAAAAGA	AGTCACTTTC	ACACTAGAGC	AAAAAGACTA	CAAGGGACTT	GCTAAAGACA	18120
TCTCAGAAAA	TGGAAAACCTT	TTAGTTCAAT	GTGATAACGG	AAAAGAAATC	TGGCTAAATA	18180
GTGGCGAAAT	TTCTCTCAAT	AGTTGGAAGT	AAAATAACAC	AATTATAATA	TAACAGATAT	18240
AAAAATAACT	TCAGATTAGT	AATTCAATTA	AGTTTACGG	ATCTGAAGTT	TTATTGGCTC	18300
TAAAAATAAA	AAAGAGAGTT	ACAGACTCTC	ATTAAAACGG	AGAATAAGGG	ATTCGAACCC	18360
TTGCGCCAGT	TACCCGACCT	AACGATTTAG	CAAACCGTCC	TCTTCAGCCT	CTTGAGTAAT	18420
TCTCCAATTA	ATGGGCACGA	GTGGACTCGA	ACCACCGACC	TCACGCTTAT	CAGGCGTGCG	18480
CTCTAACAC	CTGAGCTACG	CGCCAAGTT	AAAAAACTTG	GTAATTGAA	CAAAGTTCAA	18540
AGCGGGTGAC	GAGAATCGAA	CTCGCGACAA	CAGCTGGAA	GGCTGTAGTT	TTACCACTAA	18600
ACTACACCCG	CATAAATACT	ATCAATAAAA	TGGCGCGAGA	CGGAATCGAA	CCGCCGACAC	18660
ATGGGAGCTTC	AATCCATTGC	TCTACCAACT	GAGCTACCGA	GCCTTATTGC	GGGAGCAGGA	18720

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TTTGAACCTA CGACCTTCGG GTTATGAGCC CGACGAGCTA CCGAGCTGCT CCATCCCGCG	18780
TTAATAATAT AAAAGGAGGA TGTGGGATTG GAACCCACGC ACGCTTTAC ACGCCTGACG	18840
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AAGGTCCGAC AAGATCATTA TAGCCGCAGA GGGGATCGAA CCCCCGACCT CCCGGGTATG	19020
AACCGGACGC TCTAGCCAGC TGAGCTACAC CGCCATGAAT CGGAAAGACA GGATTCGAAC	19080
CTGCGACACC TTGGTCCCAA ACCAAGTACT CTACCAAGCT GAGCTACTTC CCGAGTTAAA	19140
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TCTAACCGAA CGACGGGATT CGAACCCCGCG ACCCCCACCT TGGCAAGGTG GTGTTCTACC	19380
ACTGAACTAC GTTCGCACTG TTTTCTTCTA TCTAAAAATG CCGGCTACAT GACTTGAACA	19440
CGCGACCCTC TGATTACAAA TCAGATGCTC TACCAACTGA GCTAACGCCG CTCATTGTT	19500
ATATCTTAAT GCGGGTTAAG GGACTTGAAAC CCCCACGCCG TTAAGGCCA GATCCTAAAT	19560
CTGGTGCCTC TGCCAATTCC GCCAAACCCG CATATATGAC CCGTACTGGG CTCGAACCAG	19620
TGACCCATTG ATTAAAAGTC AATTGCTCTA CCAACTGAGC TAACGAGTCT AAAATAACTT	19680
CGCTTACCTT AAACGGTCCG ACGBAATCGA CCCGGTAC	19718

(2) INFORMATION FOR SEQ ID NO: 100:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4117 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 100:

CCGTGGAAAA GTCTGGATAG TGAATGGTCT TCACACAATG ACCTGAAAGA AGCCTGAGAA	60
TAATTATGGA GAGTAGCATT CTGAGAGGTG TTAGCAGAAC CATATGACAG AGCTGTTGA	120
AGAGGGAAATA TTGAGGAGAA AAATCCTGAG CCTACCAGTT GGAGTTGGAA AGAGCTGACT	180
GTTAGATCAT GGTATTATCC ACACAAACCTG TGGATAACTT TGTGAATAAG AGAAGTTGCT	240
AAAGAAGGAG ATATATAACG ATGAAGAAAA TCAAACCGCA TGGACCGTTA CCAAGTCAGA	300
CTCAGCTAGC TTATCTGGGA GATGAACCTAG CAGCTTTAT CCACCTCGGT CCTAATACCT	360
TTTATGACCA AGAATGGGGG ACTGGACAGG AGGATCCTGA GCGCTTTAAC CCGAGTCAGT	420

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TGGATGCGCG	TGAGTGGTT	CGTGTGCTCA	AGGAAACGGG	CTTCAAAAAG	TTGATTTGG	480
TGGTCAAGCA	CCACGATGGC	TTTGTCTTT	ATCCGACAGC	TCACACAGAT	TATTGGTTA	540
AGGTCACTCC	TTGGAGGAGA	GGAAAGGGCG	ACTTGCTCCT	TGAAGTATCC	CAAGCTGCCA	600
CAGAGTTGA	TATGGATATG	GGGGTCTACC	TGTCACCGTG	GGATGCCCAT	AGTCCCCTCT	660
ATCATGTGGA	CCGAGAACG	GACTACAATG	CCTATTATCT	GGCTCAGTTG	AAGGAAATCT	720
TATCAAATCC	TAACATATGG	AATGCTGGTA	AGTCGCTGA	GGTTGGATG	GATGGTGCCA	780
GAGGAGAGGG	CGCGAAAAG	GTAAATTATG	AATTTGAAAA	ATGGTTGAA	ACCATTCGTG	840
ACCTGCAGGG	CGATTGCTG	ATTTTTCAA	CAGAAGGCAC	CAGTATCCGC	TGGATTGGCA	900
ATGAACGAGG	GTATGCAGGT	GATCCACTGT	GGCAAAAGGT	GAATCCTGAT	AAACTAGGAA	960
CAGAACGAGA	GCTGAACTAT	CTTCAGCACG	GGGATCCCTC	GGGCACGATT	TTTTCAATCG	1020
GAGAGGCAGA	TGTTTCCATC	CGTCCAGGCT	GGTTCTACCA	TGAGGATCAG	GATCCTAACT	1080
CTCTCGAGGA	GTTGGTCGAA	ATCTACTTTC	ACTCAGTAGG	GCGAGGAAC	CCACTCTTGC	1140
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AATTGCGAC	CTATCGAAC	GAGCTCTATA	AAGAAGATTT	GGCTCTGGGA	GCTGAGGTAT	1260
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GCTCTGGGC	AAGCGATGCA	GACTGCCA	TCCAGTTAGA	ACTCGACTTA	GGTTCTCCTA	1380
AAACTTTGAG	TGTAATTGAG	TTAACAGAAC	ATTGAAGCT	AGGGCAACGA	ATCGCTGCTT	1440
TTCATGTGCA	AGTAGAGGTG	GATGGTGTCT	GGCAGGAGTT	TGGTTCGGGT	CATACTGTTG	1500
GTTACAAACG	TCTCTTACGA	GGAGCAGTTG	TTGAGGCACA	GAAGATACGT	GTAGTCATTA	1560
CAGAACATACA	GGCTTTGCCT	TTGTTGACCA	AGATTTCCCT	TTATAAAACT	CCTGGATTAT	1620
CAAAAGGAGA	AGTTGTTCA	GAACATAGCAT	TTGCAGAAAA	AAGCCTAGCT	GTGGCAAAGG	1680
GAGAAAATGC	CTATTTACA	GTAAAGCGCA	GAGAATGTAG	TGGTCCTTTA	GAAGCTAAGA	1740
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TTGCGTTCA	AACTGGTGAG	ACTGAAAAAA	GTCTGACGCT	ACCAACCTTG	TATTCGCGAG	1860
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TTCAAGTCCA	AGTTTCATAA	AAGAAGAAC	TTTGCAGCAG	GCAAAGGTTC	TTTTGGTTAT	1980
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TGTTGAATAG	TTGATACGAG	TGTTTGCTC	AGTCGGCATT	CTTTGACAAA	GTTAAAATGG	2100
TTGTGGTTTT	GTTTAGTATG	GATATCCAGC	CATTTATCTT	CTTTAGCGAG	GTAGACTCGT	2160

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AGATGGTCAA AGAGAGGGAT TCCGAGGTCA TAGCTGGTT TTCCCTGGACA GGTTGGATAA	2220
AATCCGAGAG CTGACCAGAT GTACCAAGCA GAGAGACTAC CATTGTCTTC ATCTCCAGGA	2280
TAGGCTTCCC AACTTGGGTG AAAAGCTTTC TGACGGAGCG TCTTGATAAG AAGGGCAGTG	2340
TAGTCAGGGT AATCGCTGTA ACGGAAGAGA TAAGGAATGT GGAAACTAGG CTGGTTGGAA	2400
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TAGCCTGTTG TTTCAGGAG GGGAGCATCT TGACAGGCTT TCAAAAGATA GTTGCTAAAG	2520
GTTTCTTTTC CACCCATCAG TTGGATTAAG CCAGGGATGT CGTGGAGAAC GCCTAAAGTA	2580
GCTTGAATGG CAGAGCATTG AGCGTAGTCT CGCCCCAAC TATAAGGAGA GAAGTCAGGG	2640
TGAAAGTTTC CTTGATGTGTC TCGTGCTCGC ATGTAACCTG TCTCAGCGTC AAATAGCTGG	2700
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GCACAGCTGG CGATACAAAA GTCACTATAG GCATAGTCTA GAGTATGGCT AACACTTCG	2820
TGGTGGTCGG TAGAGAGGTA ACCTAGTTCT TGGTATTGGG CTAGTCCGTG GCGGCCATTG	2880
ATGCCGAGAG GGTCGGCTT GCTGGCTTT TCGAGCATGG CTTGGAAGAG TTCTCCTTCT	2940
AGGTGGGGGG TCATGTCCCTT GCAGGGCTA TCTGCGATAA TACCGTCTAA AAGTGTACCT	3000
GGCATCATAAC CCCGTTCATC TGGAGCCAGC CATTGGAA GGAAACCAGT ATCGCGGTAG	3060
CTATTGAGGA AACCTTCTAA AAAGCGTTGA TAGTGCTCCG GTATGATAAG GGCAAAGAGG	3120
GGGAAGGTGG TGCAGGAGGT ATCCAGAAA CCATTGTTGC TAAAGAGGAC ACCAGGCTTG	3180
ACAGTACCAAG TAGCCAGATC CATGTGGATG GCTTGCCCTG ATTCAATTAACT CTCATAAAAAA	3240
GTCTGTGGGA AGAGGAAGAG TCTGTAGAGG CAGTGGTCAA AGAAGGTTG GTCAGCCTCT	3300
CCTGTCTCTA TAATGTCAAA ACGATGGAGG AGATTTCCC AATCCACTTG GGCACATTGAT	3360
TTACAGCTAT CAAAATCTTC TTGAGGTAGA TTGATTAGAG CTTGAGAAGG AGAGATGAAA	3420
GAAGTGGCTA GTTGCATCTC GGTTGACTA CTTGCTAAGT CAATTGCGCA GTCTCCAGCT	3480
TCTTGGCTGA TAGCAAGAAT ATCCGTGTTA ATTTGCAGGG CAGTGAACAT CGTTAGCGAA	3540
TTTTGGTTAG TTTCAGTTTT ACCTTCTTGT CGCAGGGCAA GAGTCCGCTT ATCTACTTGC	3600
TCTACTGTCA GTTCATCTGC TGCGTGAAGA TAGAGGGAGA GGGCTTGCC TTGCTTTGA	3660
TTCAAACGAA TAGAACGACC ATAGCAAGTC GGTGTGAGCT GGGTTCAAT CTGATAACGC	3720
AGAGAAAAGA GCTTCAAATA GTGAGGCTGG AAGCAAGCTT TATCTATATC ATAAGAAGAC	3780
TGGCGGTGAA AGAGGCTGTC TCCCCCAGT TGACTGGTGA CAGGTGTCAG AAGGAGCCAA	3840
GAGTAGTCCC CAATCCAAGG ACTGGGCTGG TGAGTTAACATC GAATCCCCTG AAAGATAGGC	3900
AGATGTGGAT CAAAAAACCA AGATCCATCC TGGTCACTGG TCTGGGGCAC AAAGTAATTG	3960

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ATCCCAAAAG GCACGCCGT GTATGGCAGG GTATTTCCC GAGAAAAGGC ATGCTTGTG	4020
GTAGTTCCA AACGGGTATC GATGGTATCA AGTAGTGGTT TCATAGTCCT TCCTTTAGCT	4080
GTTTTCTAC ATTATATCAC TAATAGAGGG CCTTTAG	4117

(2) INFORMATION FOR SEQ ID NO: 101:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2727 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 101:

CTGGTTCAAT TATTATTCAC TCTAAAGTAGT CATATGTTCT TTATTTATGT GAGTTTTTAC	60
CTTTTAAAGG ATCTTGTTAG ATGGGAGAAG GTTTTAAAAG TGACAGATGA TAATACAAGA	120
AAAGTTCGTT TATTAGTAGC CTTTTTAGC ATTGTATAG GCTACATCCT GAGTTCTTC	180
TTTATTAGCC TGTATCATTT GTGGCAAGAA GCGCTTAGAG GATTATTATG AAATCAAGAG	240
TAAAGGAAAC GAGTATGGAT AAAATTGTGG TTCAAGGTGG CGATAATCGT CTGGTAGGAA	300
GCGTGACGAT CGAGGGAGCA AAAATGCAG TCTTACCCCT GTTGGCAGCG ACTATTCTAG	360
CAAGTGAAGG AAAGACCGTC TTGCAGAATG TTCCGATTTT GTCGGATGTC TTTATTATGA	420
ATCAGGTAGT TGGTGGTTTG AATGCCAAGG TTGACTTTGA TGAGGAAGCT CATCTGTCA	480
AGGTGGATGC TACTGGCGAC ATCACTGAGG AAGCCCCTTA CAAGTATGTC AGCAAGATGC	540
GCGCCTCCAT CGTTGTATTA GGGCCAATCC TTGCCCCGTGT GGGTCATGCC AAGGTATCCA	600
TGCCAGGTGG TTGTACGATT GGTAGCCGTC CTATTGATCT TCATTTGAAA GGTCTGGAAG	660
CTATGGGGGT TAAGATTAGT CAGACAGCTG GTTACATCGA AGCCAAGGCA GAACGCTTGC	720
ATGGTGCTCA TATCTATATG GACTTTCCAA GTGTTGGTGC AACGCAGAAC TTGATGATGG	780
CAGCGACTCT GGCTGATGGG GTGACAGTGA TTGAGAATGC TGCGCGTGAG CCTGAGATTG	840
TTGACTTAGC CATTCTCCTT AATGAAATGG GAGCCAAGGT CAAAGGTGCT GGTACAGAGA	900
CTATAACCAT TACTGGTGT GAGAAACTTC ATGGTACGAC TCACAATGTA GTCCAAGACC	960
GTATCGAACG AGGAACCTTT ATGGTAGCTG CTGCCATGAC TGGTGGTGT GTCTTGATTC	1020
GAGACGCTGT CTGGGAGCAC AACCGTCCCT TGATTGCCAA GTTACTTGAA ATGGGTGTTG	1080
AAGTAATTGA AGAAGACGAA GGAATTCGTG TTCGTTCTCA ACTAGAAAAT CTAAAAGCTG	1140
TTCATGTGAA AACCTTGCCC CACCCAGGAT TTCCAACAGA TATGCAGGCT CAATTTACAG	1200

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CCTTGATGAC AGTTGCCAAA GGCGAATCAA CCATGGTGGGA GACAGTTTC GAAAATCGTT	1260
TCCAACACCT AGAAGAGATG CGCCGCATGG GCTTGCATTG TGAGATTATC CGTGATACAG	1320
CTCGTATTGT TGGTGGACAG CCTTTGCAGG GAGCAGAAAGT TCTTTCAACT GACCTTCGTG	1380
CCAGTGCAGGC CTTGATTTG ACAGGTTGG TAGCACAGGG AGAAAATGTG GTGGTAAAT	1440
TGGTTCACTT GGATAGAGGT TACTACGGTT TCCATGAGAA GTTGGCGAG CTAGGTGCTA	1500
AGATTCAAGCG GATTGAGGCA AGTGTGAGAAG ATGAATAAGA AATCAAGCTA CGTAGTCAG	1560
CGTTTACTTT TAGTCATCAT AGTACTGATT TTAGGTACTC TGGCTCTAGG AATCGGTTA	1620
ATGGTAGGTT ATGGAATCTT GGGCAAGGGT CAAGATCCAT GGGCTATCCT GTCTCCAGCA	1680
AAATGGCAGG AATTGATTCA TAAATTTACA CGAAATTAGG CTGGAGAACCG AGCCTTTTC	1740
TAAAGATAAG GAGAAATATG AACAAAAAAA CAAGACAGAC ACTAATCGGA CTGCTAGTGT	1800
TATTGCTTT GTCTACAGGG AGCTATTATA TCAAGCAGAT GCCGTCGGCA CCTAATAGTC	1860
CCAAAACCAA TCTTAGTCAG AAAAAACAAG CGTCTGAAGC TCCTAGTCAA GCATTGGCAG	1920
AGAGTGTCTT AACAGACGCA GTCAAGAGTC AAATAAAGGG GAGTCTGGAG TGGAATGGCT	1980
CAGGTGCTT TATCGTCAAT GGTAAATAAA CAAATCTAGA TGCCAAGGTT TCAAGTAAGC	2040
CCTACGCTGA CAATAAAACA AAGACAGTGG GCAAGGAAAC TGTTCCAACC GTAGCTAATG	2100
CCCTCTTGTC TAAGGCCACT CGTCAGTACA AGAATCGTAA AGAAAATGGG AATGGTTCAA	2160
CTTCTTGGAC TCCTCCAGGT TGGCATCAGG TCAAGAATCT AAAGGGCTCT TATAACCATG	2220
CAGTCGATAG AGGTCAATTG TTAGGCTATG CCTTAATCGG TGTTTGGAT GGTTTGATG	2280
CCTCAACAAAG CAATCCTAAA AACATTGCTG TTCAGACAGC CTGGGCAAAT CAGGCACAAG	2340
CCGAGTATTC GACTGGCAA AACTACTATG AAAGCAAGGT GCGTAAAGCC TTGGACCAAA	2400
ACAAGCGTGT CCGTTACCGT GTAACCCTTT ACTACGCTTC AAACGAGGAT TTAGTTCCCT	2460
CAGCTTCACA GATTGAAGCC AAGTCTTCGG ATGGAGAATT GGAATTCAAT GTTCTAGTTC	2520
CCAATGTTCA AAAGGGACTT CAACTGGATT ACCGAACGG AGAAGTAAC GTAACTCAGT	2580
AAAAGATACG CCTACACTCC TATGTCACTT ATGGATGTAG GAGTTCTTT TACTAGTTA	2640
ACCAGGACTA AGACAGGTAC TAAGACAAAA TAGCAACTTC TAAAACTAAC TTCCAGTTT	2700
GGGAGAGAGA TGGAAGTTAC TTTGAGA	2727

(2) INFORMATION FOR SEQ ID NO: 102:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5717 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 102:

TTTTTTGTAG ATTTAAGTGG GGTGCAATT CTAACAAATA AAAAACAAATT TTTGAAAATT	60
ATGTTAGCAG GAATTGCTTC AAATTCGATT TTATCACTTA CAGGTTACT TGTTTATTG	120
TTCACACATCGT ATAAATTGCT TGGACTCTTA TTTTTTATCA TTAACCTAGG TATGATTTT	180
ATTAATTCAA TTCCTTTTT TCAGTATGAT AGTGGTATTA TTTAAAGATA CTTGAATTCT	240
AACAATAATA ACTTGAATT TCAATATATA GTTCAACTTT TAATAGCATT TGTTATTATT	300
TATTTTCCTT TGAGTCAACT ATTACAGTTT TTGACACCCA ATATTATTGT TCGTAGTATA	360
GGAGGGGTGG TTGTTTCTAT ACTGCTTCT ATATTATATA TGATAGGAAG GACGAAATAT	420
GTTCTACGTA AATAGTTATG TTTTGCTTA TAAAAAAGAA GGTATAATGT ATTTACGTGG	480
TCGGAGTATG CGGGAAATAG CTATAGAACCC TCAAATTTCG CAAGAATTAA TCAACGATCT	540
ATTTAATAGT TGTAAGAAC TATTAGAGAT AGAAGAAGTA TTAGGCAGTA AACTAACATT	600
TGAACTATAA ATGAACAAAT TTTAATTTCG GATGAGATAG ATATTGATAG TAGATATTCT	660
AGAACTAAAG GTTACTATTG GTTATTCTAT AATGAAGAGT ATAATAAAAT ACAGAATAAA	720
ACAGTATTAG TATTAGGAGC AGGAGTCTTA GGATGTTATA TATCTCTAAG TCTAAGTATG	780
TATGGAGTGA GGAAACTTAT TGTCGCTGAT TACGATATAA TAGAACCATC AAATTTAAAT	840
AGGCAAATTC TTTATACAGA GTCGGATGTT GGTAAGGAGA AGATTAATGT TCTTCTGAA	900
AAAATACACA AGTATAATTG AGATGTTCAAG GTAGTACCTA TTTCTATTAA AGTTTCTTCA	960
GTAGAAGAAT TAGAAAAAAT TGTTGCGGAA TATGGGAGTA TAGATTCTAT CGTAAAGCA	1020
ATTGATACGC CCATTGATAT TATAAAATT GTCAATCAAT TTGCTGTATC GCATAAGATA	1080
TCCTACATAT CAGGAGGTT TAATGGATGC TATCTTATTA TTGATAATAT ATATATCCCT	1140
ACCATCGGTT CTTGCTTGG TTGTCGGAAT ATAAACAAAG ATATAAATAA GTACACTTAA	1200
TCTGATAAGA CAAAGTGGCC GACTACACCA GAGATGCCTG CTATTTGGG AGGGATAATG	1260
ACTAATTAA TAATTAAAT ATTTCTGGGA TGTTATAATG AAATCCTAAT AGATAACGCT	1320
TACGTTATA ATATGAGAAA TCATGCTCTA AGTCAAGAAA AATATGTTCT GGAAAACGGA	1380
GAATGTCCAA TTTGTAAAAA AATAATAAAAG TGAAAGATAA CAATATTAGA GCGAAAACAT	1440
TTATTCGTTCA AGTTGTTTT TGCTTATTAT CAGGAGGAGT AGCTTTTTA TCTGCTATTG	1500
GGCAGTTCAC TGTTATAGAA ACACAATTAA TAGTATTGTT CTTGGGTATT ATTTTGCTA	1560
TATATTATGC TTACTACAAT AAAAATATTG AAACATCATT GGAAAATATA GTATGGCTTT	1620

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TTTCATCGTT	TGAGATTTA	TTTTGCTTG	TTAATTAG	AACATTATT	CAGTTACCAG	1680
TGGATATTT	TATTGGTATG	ATAATATT	TAATGCTGTG	GATATTATT	ATGTTAGGTA	1740
TAGTGTGTCT	TAGTTATTAT	ATAACTTAT	TATTTAGCAA	GGAGGCTTAG	TATGTTAAA	1800
AAAATAGGTA	TAATGAGCAT	TTGCATATAT	ATAATTATT	TATACTGCTT	GAGAATGTAT	1860
CGTATTATCA	ATAATATTGA	AACAATCTTG	CTAACGGTTA	TATGCTTAAT	GTTATTGTTT	1920
TTTTAAGAC	GTTTATTGTA	AAAGATAAG	AAATAGATG	TTAAGTAAAA	ATGTAGAATA	1980
TAAAGGAGGT	GCAATGAGTA	TGATTGAAGT	TAGCCATTAA	TCAAAAAGTT	TTGGTGATAA	2040
AATAGCTTTA	AATAATATAA	GCTTCACTGT	TAAAGAAGGT	TAGATTTTG	GATTTTAGA	2100
ACCATCTGGT	TCTGGAAAGA	CCACAACGAT	TAATATTCTG	ACTGGGCAGT	TCCTGCCGA	2160
TAAAGGACAA	TCTATTATTT	TGGGACAAAA	ATCTCAAAAT	TTAACAAAGCG	GTGAATTAAA	2220
GAGAATTGGA	TTGGTTAGCG	ATACAAGTGG	ATTTTATGAG	AAAATGTCTC	TGTATAACAA	2280
TCTTCTTTT	TATAGTAAAT	TTTATAATAT	TAGTAAATCA	CGTGTGATA	ATTTGTTAAA	2340
GCGAGTAGGA	TTATATGATA	GTCGCAAGAT	GGTAGCAGGA	AAATTATCCA	CTGGAATGAG	2400
GCAACGAATG	CTTTTAGCAC	GAGCTCTTAT	CAACAACCCC	GCTGTACTCT	TTCTGGATGA	2460
ACCGACCTCA	GGTCTAGATC	CCACAACCTTC	TCGAACAATT	CATGAGTTAA	TTTTAGAATT	2520
GAAAACAGCA	GGGACAACGA	TTTTCTAAC	GAETCATGAT	ATGAATGAAG	CAACTCTTTT	2580
ATGTGATTAT	GTTGCCTTAT	AAATAAAGG	GAAATTAGTT	GAGCAAGGAG	CTCCTCTGA	2640
ACTCATTCAA	AGATATAATA	AAGATAAAAA	GATTAAGGTT	ACAGATTATA	ATGGGAATCA	2700
GATAACTTT	GATTTACAT	CACTAGAACCA	GGTATCTCAG	ACTGATCTGG	AAAATATT	2760
TTCAATTCA	TCATGTGAGC	CTACTTTAGA	AGATTTTTT	ATCACATTAA	CAGGAGGAAA	2820
GCTAAATGCT	AAACGGTTT	CTGGCTTGG	TATGGTTGCG	TTGTCAAATC	ATCCTTCCA	2880
ATAAGAGTAT	TTTATTGCAA	GTGTTAGTGC	CTTTGCTTT	CACATATT	TATAAATATC	2940
TTATGGAAAC	ACAGGGGAAG	GTCAACGATC	AACAGGCATT	AGTTCTTTG	ATGATGTGTT	3000
TACCTTTTTC	TTTTCTTTG	GCTGTTGGAA	GTCCTATAAC	TATTATCTTG	TCTGAAGAAA	3060
AAGAAAAGTA	CAATTACAA	ACTCTCTGT	TGAGTGGTGT	AAAGGCTCC	GAATACATT	3120
TATCAACTAT	GTTCCTCCT	TTTTGCTAA	CTTTGTGAT	TATGGGAACT	ACTCCTTTA	3180
TTTTAGGAGT	TACAATTGTA	CATACTTTA	ATTATATTAC	AATCGTTCTT	CTAACCTCTT	3240
TATCCATCAT	TTTATTCTAT	TTATTGATAG	GTGTTAACCGC	GAAGAGCCAA	GTAGTAGCTC	3300
AGGTTATCAG	TCTTCCTGCT	ATGATTTAG	TTGCTTTCTT	ACCGATGCTA	TCTGGTTGG	3360
ATAAGACAGT	TGCGAAGATA	ACAGATTATA	GTGTTATGGG	ACTATTACT	AAGTTTTCA	3420

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CAAAATGGGA GGAATTTCAGGAAATAAAA CTCTAATTCC TAATCTAACAA CTACTTATTT	3480
GGATTGTTCT TCTATTAACCTTAATTACGA TAACTATTAG GAAAAAGAAA ATTTCTTAAT	3540
TGAGTTATTT TAATGATTAT AAACACAAAGT GGGAAAGGAAA AAATGAACCTG ATCTTTTGAA	3600
CAGCAATTCT ACAGAACATGT CTTATTGCTA TATTTTGATT TGAGTGTACG AAAAAAGAAA	3660
AATAACAATA GTGCTCATACT TAATTGCAGA AGTTTGCGGT GATAAGATAA CTGATAAAATT	3720
GCAATAAAAAA ATGCAACATT TTAAATCTC CTCTATAAGT GCTTCAAAAA GTGCTTCAAA	3780
ACCTGTCTTG TAATCCAAGT ATTTTGCGGT ACGGTGATTA ATAAGCTAGC AAAGCATCAT	3840
TAAGGATTTT TTCGGTAATT GTGCCAAAT CGGTTTAAGA AAATACTCAC GAAGAAGTCC	3900
ATTCGCATTTC TCATTACTTC CCCTTGCCA AGATGAATAG GCATCCGCAA AATAAACAG	3960
AATTCCCATT TGTTCAATTA AAGGGTAACA AGCAAACTCT TTTCTCTGT CCGAAGTGAA	4020
AGTCTTTAAC TATTCTTTG GAAAGAGTCT TGTGAGGTGT TCAATAGCAG TCAACATGGA	4080
TTTAGCTGTT TTACTTGAC AAGTGTAGT AGAAATAATA GAATAGTAA AACCTTTAA	4140
AGCAGTCCAG AGAGGCAGCT AAGGTTAGAC GGTGAAAGGG TGGAGACTAC CCATTTCG	4200
TGGAACCTTG CTGTTGGCAG GTTCCCTTTTC TCGTGGCTTC TGTTGGCCAG ACTCTCTCAC	4260
TAGTAAAGGT AAAAGGAGAA ACCTATGCGA GAACATCGTC CAATCATTGC TCTTGATTTT	4320
CCTAGTTTG AGGCGGTCAA GGAAATTTTA GCTCTTTCC CAGCAGAAGA AAGCCTTTAT	4380
CTCAAGGTAG GGATGGAGCT TTATTACGCA GCGGGGCTG AGATTGTGTC CTACTTAA	4440
GGTTTGGGTC ATAGTGTCTT TTTGGATCTC AAACTTCATG ACATTCTAA TACAGTCAAG	4500
TCAGCCATGA AGATCTGTGTC TCAGCTTGGT GTCGATATGA CTAATGTCCA TGCGGCTGGT	4560
GGTGTAGAGA TGATGAAGGC GGCGCGTGAA GGTCTTGGGA GTCAAGCCAA ATTGATCGCT	4620
GTAACTCAGC TCACATCAAC GTCAGAACAGT CAGATGCAGG AGTTTCAAA TATCCAAACC	4680
AGTCTGCAAG AGTCTGTGAT TCACATATGCC AAGAAGACAG CTGAAGCTGG CTTGGATGGT	4740
GTTGTTGCT CGGCTCAGGA AGTACAAGTC ATCAAGCAGG CTACCAATCC AGATTTTATC	4800
TGTCTGACAC CAGGGATTGTC TCCAGCTGGT GTTGCAGTTG GAGATCAAAAC ACGAGTCATG	4860
ACACCTGCTG ATGCCTATCA AATCGGCAGT GACTATATCG TAGTGGGACG TCCCATTACC	4920
CAAGCTGAGG ATCCTGTTGC AGCTTATCAT GCCATCAAGG ATGAATGGAC ACAGGACTGG	4980
AATTAAAGAA CTAGATTAGA AAAATAAAAG GAGAATACCA TGACACTTGC TAAAGATATC	5040
GCTAGGCCACC TCTTGAAAAT CCAAGCCGTT TACCTCAAAC CAGAGGAACC CTTCACTTGG	5100
GCATCTGGTA TCAAGTCACC GATTACACT GATAATCGTG TGACACTAGC CTATCCAGAA	5160

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ACTCGTACCC TAATTGAAAA TGGTTTGTC GAGCTATCA AAGAACCTT TCCTGAAGTA	5220
GAAGTGATTG CAGGAACCTGC AACAGCAGGG ATTCCACACG GAGCCATTAT TGCTGATAAG	5280
ATGGACTTGC CTTTGCCTA CATCCGTAGT AAACCAAAAG ACCACGGAGC TGGTAATCAA	5340
ATCGAAGGTC GCGTAGCTCA AGGTAAAAA ATGGTAGTGG TTGAAGACCT TATTCAACG	5400
GGTGGTTCAG TTCTTGAGC TGTAGCAGCA GCCAAGCGAG AAGGAGCAGA TGTACTTGGA	5460
GTTGTAGCGA TTTTCAGCTA CCAATTGCCA AAAGCAGATA AGAACCTTGC AGATGCTGGT	5520
GTTAAACTTG TGACGCTTTC AAACATAGC GAGCTTATCC ATCTAGCCC AGAAGAAGGT	5580
TACATCACGC CAGAGGCCT TGATCTCTA AAACGCTTTA AAGAAGACCA AGAAAATTGG	5640
CAAGAAGGTT AGGTCAGTAA GATAAAGAGA GACGAGGCTA CCGAGTCTCT TTTACCATTT	5700
TATTTAAAAT ATGACAG	5717

(2) INFORMATION FOR SEQ ID NO: 103:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5558 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 103:

CCTGGACTTT CTAAAATGAA ATCTTGCAC CTGGATCAAG CCCTTCATGA GCATTTTCA	60
GAAGAAGAAT TAGCTGGTCA CTTTCATGTC CTTCTATGGA CTTTTTTTAC AATGGCATTG	120
CTATCACACC CAATACCTAT CTAAGCGCCT GTTTCGTAAA CTTTATTGCA GCTCTTCCTC	180
TAAATTTCTT AATTGTTGAA CCAATTGCCA GTTTTATACT AAGTTCTTT CAGAAACCAT	240
TTACTGGGGA AGAAGTTGAA GATTTCAAG ATGATGATGA AATCCCAACT ATTATCTAAG	300
CCAGTTCTGT AAAACTACTAA TATTGAAAT CCACTTCCTT TTAGGGTGCA ATGGTTATAA	360
ATGAATTTT GAGAGGATCA GAATGAAAAA ACTAGCAACC CTTCTTTAC TGTCTACTGT	420
AGCCCTAGCT GGGTGTAGCA GCGTCCAACG CAGTCTGCGT GGTGATGATT ATGTTGATTC	480
CAGTCTTGCT GCTGAAGAAA GTTCCAAAGT AGCTGCCAA TCTGCCAAGG AGTAAACGA	540
TGCTTTAACAA AACGAAAACG CCAATTCCC ACAACTATCT AAGGAAGTTG CTGAAGATGA	600
AGCCGAAGTG ATTTTCCACA CAAGCCAAGG TGATATTGCG ATTAAACTCT TCCCTAAACT	660
CGCTCCTCTA GCGGTTGAAA ATTCCTCAC TCACGCCAA GAAGGCTACT ATAACGGTAT	720
TACCTTCCAC CGTGTACATCG ATGGCTTAT GGTCCAAACT GGAGATCCAA AAGGGGACGG	780
TACAGGTGGT CAGTCCATCT GGCATGACAA GGATAAGACT AAAGACAAAG GAACTGGTTT	840

771

CAAGAACGAG ATTACTCCTT ATTTGTATAA CATCCGTGGT GCTCTTGCTA TGGCTAATAC	900
TGGTCAACCA AACACCAATG GCAGGCCAGTT CTTCATCAAC CAAAACCTCA CAGATACCTC	960
TTCTAAACTC CCTACAAGCA AGTATCCACA GAAAATTATT GAAGCCTACA AAGAAGGTGG	1020
AAACCCTAGT CTAGATGGCA AACACCCAGT CTTTGGTCAA GTGATTGACG GTATGGATGT	1080
TGTGGATAAG ATTGCTAAGG CCGAAAAAGA TGAAAAGAC AAGCCAACTA CTGCTATCAC	1140
AATCGACAGC ATCGAAGTGG TGAAAGACTA CGATTTTAAA TCTTAAAAC CAAAAAAATA	1200
CAGTATCCAC ATTGGTACT GTATTTCTTT TACTCTCATT CTTAAGTTAA ATTATTAAGA	1260
TCCCATATTG GGTCTATCCA GCCTTCATAA AAGTCTGGCT CGTGGCAGAC CATAAGGATA	1320
GATCCCCAT ATTCTTGAG AGCGCGTTG AGCTCATCCT TTGCATCCAC ATCCAAATGG	1380
TTGGTCGGCT CGTCCAGCAC TAAAACGTTG TTTTCACGAT TCATCAAGAG ACAGAAACGA	1440
ACCTTGGCTT GCTCTCCCC TGATAAATCT TGAATCTGGC TTTCAATATG TTTGGTTGTC	1500
AAACCACAAAC GGGCAAGGGC TGACCGGACT TCTGCTTGAT TAAGGGCAGG AAAGGCATTC	1560
CAGACAGCTT CAAGAGGAGT TTGGCGATTA CCGCCTTCTA CTTCTGCTC AAAATAACCA	1620
AGTTCTAAAT AATCTCCACG CTCCACTTCC CCAGCGATTG GCGAGATAAT GCCCAAGAGA	1680
CTCTTCAAGA GAGTTGTTT TCCAATACCA TTAGCACCAA TAATCGCAAC CTTTGATTG	1740
CGTTCGAAGG TAAGATTAA AGGCTTAGTA AGAGGACGGT CGTAACCAAT TTGCAAGTTC	1800
TTGGCTTGGGAGATAAAAGCG CCCTGGTGTGTA CGAGCTGGTT TGAAATCAA GGATGGTTT	1860
GGTTTCTCAC TTTGGAGTTC GATAATATCC ATCTTATCCA ATTTCTTTG ACCAGACATA	1920
GCCATATTAC GAGTTGCAAC ACGGGCTTTA TTACGAGCCA CAAAGTCCTT GAGGTCTGCA	1980
ATCTCTTCT GCTGGCGTTC GTAGGCTGCC TCTAGCTGAG ATTTCTTCAT AGCATAAAACT	2040
TCTTGGAACT GGTAGTAGTC ACCAGAGTAA CGCGTCAGCT GTTGATTTTC CACATGATAG	2100
ACAATATTAA TAACGTCATT GAGGAATGGA ATATCGTGCG AAATGAGAAC AAAGGCATTC	2160
TCATAGTTTT GGAGATAGCG CTTGAGCCAA TCAATATGCT CAGCATCCAA GTAGTTGGTC	2220
GGCTCGTCCA ACAGCAAGAT ATCAGGCTTT TCAAGGAGAA GTTTGCCAA AAGCACCTTG	2280
GTTCTTGCC CACCTGACAA AGAAGTTACA TCCGTATCCA TGCCAAAGTC CATAACACCA	2340
AGAGCAGCGC CTACTTCGTC AATCTTAGCA TCCAAGGTAT AGAAATCACG ACTCTCCAGA	2400
CGGTCTTGAA GTTCTCCTAC TTCTTCCATG AGAGCATCAA CATCCGCGCC GTCTTCAGCC	2460
ATTTTCATAT AGAGGTCAATT GATACGAGCT TCAGCTTGA AAAGCTCATC AAAAGCCGTA	2520
CGGAGAACAT CACGCACCGA CTGTCTTCA GCAAGGACAG AGTGCTGATC CAAGTAACCA	2580

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GCCGTCACAT	ATTTGGACCA	CTCAACCTTT	CCTTCATCTG	GCAGCATT	ACCAGTCACG	2640
ATACTCATAA	AGGTTGATT	TCCTTCACCA	TTGGCACCGA	CCAGGCCGAT	ATGTTCTCCC	2700
TTGAGGAGAC	GGAAGGACAC	ATCTTCAAAA	ATTGCACGGT	CACCAAAACC	GTGACTCAGA	2760
TTTTTAACCT	CTAAAATACT	CATTCTAACCTTG	TTTTTATGTA	ATCGTTTATA		2820
AAGGAGCCAA	GCCAGATAGC	CACCCAAAGT	GTTGGTCCAC	AAATCATCAA	TCTCAAAGAC	2880
GCGATTGAAA	TCAAAGAAAAA	AGTCCAAGAT	TAATTGCGTA	CACTCGATTC	CAAGACTCAC	2940
AAGAAAAACTA	AAAAGAAGGA	CCTTTTTGT	TTTCCGAAA	TTTGGAAATA	GATAAAGGAG	3000
TTGGAAAATC	AGAGGAAAAAA	ACAAGAAGAC	ATTGAGGATA	TTTTGTAAAA	AAATCCAACA	3060
TAATTGTCCA	ATGTCACTCA	CTTCGCCAG	TTTCCAGAGA	GAATTGAAAG	GAGTCAAAAG	3120
AAAAACCAGG	CGTCCAAGAT	GCTGAATACC	TGGAGTTCCC	ACTCCCACGG	TAGATTGTT	3180
TTGAGGAGTA	AAGCAAAAAC	AGACAATGCA	AATGCTATAG	AAAATGACTC	CCCAGACCAA	3240
AATATGATTA	TAAGTCTTCT	TCATCATTAA	GGATTTACCG	CTGCGACTGC	CTTCTGGCGG	3300
TCACGTTCA	TTGTGTTAGA	GCGCAATTGT	CCACAAGCTG	CGTCAATATC	TGTACCATGC	3360
TCTTGACGAA	CCACACAGTT	GACCCCTTT	TTCTTAAGCG	TATCATAGAA	AGCCAACACG	3420
CACTCTTG	GACTACGGCT	ATATTGGTCA	TGCTCACTAA	CTGGGTTATA	AGGAATCAAG	3480
TTTACATAAG	ACAATTCTT	GATGTTCTTG	AGCAATTCA	TCAATTCCAA	GGCTTGTCT	3540
ACACCGTCGT	TGACTTCATT	AAGCATGATA	TATTCAAAGG	TTACACGACG	GTTTGTGTC	3600
TCAATGTAGT	ATTCAATAGC	AGCAAAGAGT	TTTCAATCG	GAAAGGCACG	GTTAATCTTC	3660
ATGATACTTG	AACGAAGTTC	ATTGTTAGGT	GGGTGAAGAG	ACACGGCAAG	ATTGACCTGA	3720
ACCCCTTCAT	CAGCAAAGTC	ACGAATTTA	TGAGCCAAAC	CTGAGGTTGA	AACCGTGTG	3780
TGACGAGCAC	CGATAGCCAT	TCCTTATCA	TCATTGATAG	TACGAAAGAA	ATTCAAGACA	3840
TTGTTGTAAT	TATCAAAGGG	CTCACCGATT	CCCATGACAA	CGATATGGCT	GATGCGTTCA	3900
TCCTGACCAC	GCTCATCAA	GTATTCTGA	ACCAGCATGA	TTTGCCTAC	GATTCACCG	3960
TTATTGAGGT	CACGTTGCTT	CTTAATCAA	CCAGAGGCAC	AGAAGGTACA	ACCGATATTA	4020
CAGCCGACCT	GAGTGGTCAC	ACAGACAGAT	AAACCATACT	GTTGACGCAT	GAGTACAGTC	4080
TCAATTAAACA	TACCGTCGGG	CAATTCAAAG	AGATATTGTA	CTGTACCATC	AGCAGACTCT	4140
TGCACAATAC	GTTGTTCAA	GGGATTGACC	ACAAACTGGT	CATTGAGCTT	AGCAATCAA	4200
TCCTTGGAAA	GGTTGGTCAT	TTCTTCAAAT	GACTGCACAC	GTTCACGGTA	GAGCCATTCC	4260
CAGATTGAT	CTGCACGGAA	TTTCTTTCT	CCCTGCTCCA	ATACCCATTC	CTGCATGGTT	4320
TGATGTACCA	AACTATGAAT	TGAGGGTTTC	ATTTCTTCTC	CTTATTCTCT	ACTCACTTCT	4380

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GACGAATGAC AAAATGACGT TGTCCTTGT CGCTTTCTG ACGACGTCTA TTTTCTTAT	4440
CTGCATTGCA CTTTCGTTA GTTGAGTCG GTTCTTTCC TTTCTAGAA GGTGTTCTT	4500
CTTCCGTCTT ACGCATTTTC TTGTCAAATG ATGCTCGCTT AGGGGCTTCA TTTCTAAGA	4560
CAAAATAGGC ACAACCATAA CTACAATACT CTAAAAGGT A GTCTGTAAA CGACTGATT	4620
TTTCAAGTTT TTCTCTGTT CGGTACATCCT TGTAAAAACC TCGTAGGCCA AGCTGTTCGT	4680
TGCTCCAGTC CCCCACGATA TAATCAAAC TGGTTAACAC TTCTGAAAAA CGCTGATTAA	4740
AAGTCGTAC ATCAAAGGCA TCCTTGATAT TTTCAACCAA GGAAAAAGCT ATCCCTTCCG	4800
TTTCGACCTT GTCCCCGTGT AAATGGAAC CC GGACCAGG AAACTTGTTA TAGTTGTATA	4860
ATTCAAGGTGC AATTCCTTT CGCATAGATA TCCTTTTTC ACGATTACTT AATACTTAT	4920
TCTACCATAA TTCTCTAGCAG TTAGCACGTT TCTCATAAAA ATGAAAAAAG TCTGACGATT	4980
TTGTCAGACC AGAATCTTAT AACCTAAAAA GAGAAGAAC ATTCTTCCCT CCAACTATCA	5040
TTTATTTAGCA GCTGCGTACA ATTCACTAC TTTATTCCAG TTGATTACTG AAAAGAAAGC	5100
TTTGATGTAG TCAGGACGCA CGTTGCGGT A TTTCACGTAG TAAGCATGTT CCCAACGTC	5160
CAAGCCCAAG ATTGGTTTTT TACCTTCTGA GATTGGTGTG TCTTGGTTG CTGTTGAAGT	5220
CACTTCAAGT TTCCCTTCTT TGTTGACAAC CAACCATGCC CAACCTGAAC CAAAACGAGT	5280
TGTTGCTGCT GCAGTGAAGG CTGCTTGGAA TTCTTCAAAT GAACCAAATG TTGCATCGAT	5340
TGCTGCTGCC AGTTCTGCTG AAGGAGCTGT TTTCTCGGG A GTCATCAAATT CCCAGAAAAG	5400
AGCGTGGTTC AAGTGTCCGC CACCATTGTT GATAAGTGCT TGACGGATAT CAGCTGGAT	5460
AGATTCTACA TCAGCAAGCA AGGCTTCAAG GTCTTCACCG ATTTCAGGGT GTTTTCTAA	5520
AGCTGCATTG GCATTGTTGA CATAAGTTG ATGGTGT	5558

(2) INFORMATION FOR SEQ ID NO: 104:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6735 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 104:

GGAATTGTAA ATATCATATT GTTTTGCAC CAAATATCG TCGTCAAATC ATTTATGGCA	60
GATACAAAGC TAGTATCGGA AGAACATAC GTGACTTATG TGAGCGTAAG GGTGTAATAA	120
TCCATGAAGC GAATGCTTGT TCAGACCATA TTCACATGCT TATCAGTATT CCTCCGAAAC	180

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TTAGTGTTC GTCCTTATG GGCTATTAA AGGGCAAGAG CAGTTGATG ATTTTGATA	240
AGCATGCGAA TTTAAAATAC AAATATGGCA ATCGCAAGTT TTGGTGTAGA GGCTATTATG	300
TAGATACGGT AGGCCGTAAT CAGAAAGTGA TAGCTGAATA TATTCAGAAT CAATTACAAG	360
AAGACAGAGT AGCAGACCAAG CTCACGTTAT TCGAGTCAGT AGATCCGTTT ACTGGCGAAA	420
TAAATAAGAG GAAGTAACTA AGGTGCTTA GCACCTGCTC GGGAAAGTGG TGCGCGAGGA	480
AGCTATTCG GTGGGCCTT GGCCCTGGCC GGTAGAACGCG GCTTATAGCC GCAGAACAAA	540
CCACCAGTTC ACACGGTGG TTTGATTTA AAAAACTTGA TACATAAAA TAAAAGTCTA	600
TATAAAGGAT GGTAAAATTC CTGTTGTCCG ATTGGACAA TATCCTAAAT AGTTACAATA	660
TATGGTCTAT ACTTTTCTT AGGAGAAAGC TAGATGTACA GACGTTGAG AGATTTGAGG	720
GAGGATCATG ATCTGCCCA AAAGCAAATA GCTACAATAC TTTCGTTAC AAATTCAAGCT	780
TATGCCAAA TTGAACGGGG TGAGCATGCG TTGACGGCTG ATGTATTGGT TAAACTCTCA	840
GATTCTATG ACGTCAGTAC AGACTATTTA TTGGGATTAA CTGATTTCC TGATAAAATT	900
CGCTTAGAA AATAATCTCC TCAATTCAT AGAGTTGAA AATGAGTGAG ATTTTTTATT	960
TGCCCTTTGA CAACTGAATA GCCTAAAATG GTACTTTCTT CATTGTGGA GCAAATTGA	1020
ATGGCTCGCC ATGATAAGAG CGATTTAAA ATCATCAATA AAATAGAGCG ATACTTTATA	1080
TGCCATGATA CAAATGATAT ACAATGATAC TTCTGACCGT TCAGCCTGCC AACGTAAAAG	1140
AGCAGCAAGT GAAATTCTTA TGATGACTTC ATCAGTCATG CCACGTTGAA TGTGTGAGTT	1200
TGTTAGATAA ACGCAATTAA TCCTCAAAG GTTCCCCGAA CCTTTGAGT TCTACAGACG	1260
CATCACGTGG AGTGTGTAAG CTTGTTGCTA AAAGCGTAAA AACCTGGAA CGAAAGGAAT	1320
AATAGACTTT CTGCGAAACA AAAATATAAT ACAATAAAAC TATGAATGAT GAAGCAAGTA	1380
AACAATTGAG CGATAGCCGT TTCAAGATCC TTGTAGGTGT TCAGCGCACG ACTTTGAAG	1440
AGATGTTAGC TGTGTTAAAA ACAGCTTATC AACGTAAACG CGCAAAGGT GGACGAAAAA	1500
GCAAATTAAG CCTAGACGAT CTCCTTATGG TAACTATTCA ATACATGCGA GAATAGAGCA	1560
CTTATGAACA AATTGCGGCT GATTTGGCA TTCACGAAAG CAACTTAATC CGTCGGAGTC	1620
AATGGGTTGA AGCAACTCTT ATTCAAAATG GTTTTACGAT TTCAAATTCT GCCTTAATTC	1680
TGTAAAACA GTAAAATTCG AAGGATTGTA AGGTAAGAGT TTTTTCTTT CTGAAAAAAT	1740
GGTATAATAG CAATCAAAAC TAGAAAATAA AACGGAATTG GGAACAGATT TGTCTGTATC	1800
CTAGTAGAGT GGTGATACTA TGAAGATTAG TAAGAGGCAC TTATTAAATT ATTCCATCTT	1860
GATTCCCTAC TTGCTTTAT CTATTTGGG CTTGATTGTC GTCTATTGCA CCACCAAGTGC	1920
TATTTAATT GAAGAAGGCA AGAGCGCCTT GCAGTTGGTT CGAAACCAAG GAATCTTTG	1980

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GATTGTTAGT TTGATACTGA TTGCCTTAAT TTATAAATTG AGACTAGATT TTTTGAGAAA	2040
TGAGCGACTA ATCATTTCAG TTATATTAAT AGAAATGCTT TTATTGTTCT TGGCTCGTT	2100
TATTGGTATT TCCGTAAACG GGGCATACGG TTGGATTCG GTTGCAGGAA TAACTATTCA	2160
GCCAGCTGAG TACTTAAAAA TCATTATTAT TTGGTATTAA GCTCACCGAT TCTCCAAACA	2220
GCAAGAAGAA ATAGCTACTT ATGATTTCA AGTTTGACT CAAAATCAAT GGCTTCCCCG	2280
TGCTTTAAAT GATTGGCGAT TCGTCTCCT AGTTCTGATT GGAAGTTGG GAATTTCccc	2340
TGATTTAGGA AATGCGACTA TTTTAGTCTT GGTTCCCTG ATTATGTATA CAGTTAGTGG	2400
AATCGCTTAT CGCTGGTTT CAACCATTCT GGCCTCGTA TCTGCCGCTT CTGCTTTGT	2460
CTTGACCAC TACAGCCTAA TCGGTGTTGA GACCTTTCA AAAATTCCAG TATTGGCTA	2520
TGTAGCCAAG CGCTTCTG CCTTTTTAA TCCCTTGCC GATCGTGTG ATGCAGGTCA	2580
CCAGTTAGCT AATTCTTATT TTGCATGGT CAATGGCGGT TGGTTGGTC TAGGTCTTGG	2640
AAACTCGATT GAAAAACGAG GTTATTTGCC AGAAGCTCAT ACAGACTTTG TCTTTCTAT	2700
CGTGATTGAA GAATTGGCT TTGTTGGTGC CAGTCTTATT TTAGCTCTCT TGTTTTCAT	2760
GATTTTGCAG ATTATCTTGG TCGGTATCCG AGCGGAGAAT CCTTTCAATG CCATGGTTGC	2820
ACTCGGTGTC GGAGGGATGA TGTTGGTCA GGTATTTGTC AATATCGGAG GGATTCGGG	2880
CTTGATTCCA TCTACAGGAG TGACTTTCCC CCTCTTATCC CAGGGTGGAA ATAGCTTCT	2940
AGTCTTATCA GTGGCAGTAG CCTTTGTCTT AAATATTGAT GCCAGTGGAA AACCGCCTAA	3000
ATTGTACCGA GAATTGGAAA ATCAACCAAT GAAACCTCTG TTGAAGTAGG ATAAAGAAAG	3060
GATAGTTAT GTCTCTCAA AAATTAGAAA ATTATAGTAA TAAAAGTGT GTGCAAGAAG	3120
AAGTCTTGAT TCTAACAGAA TTACTGGAAG ATATTACTAA AAATATGCTT GCCCCAGAGA	3180
CCTTGAAAAA AATAATACAG TTGAAAGAAT TATCAACGCA GGAAGATTAT CAAGGTCTAA	3240
ACCGTCTAGT GACTAGCTTA TCAAATGATG AAATGGTCTA TATTTCACGC TATTTCTCTA	3300
TCTTGCCCTCT TTTGATTAAT ATTCAGAGG ATGTGGATTT AGCTTATGAA ATCAATCATC	3360
AAAATAATAT TGATCAGGAC TATTTAGGTA AATTATCTAC AACGATTAAA TTGGTAGCAG	3420
AAAAGGAAAA TGCCGTTGAG ATCCTAGAAC ACTTGAATGT TGTCCCTGTT TTGACAGCCC	3480
ATCCAACACA AGTGCAACGC AAAAGTATGT TGGATTTAAC AAATCATATT CATAGTCTTT	3540
TGCGTAAATA CCGTGATGTT AAGTTGGGT TGATCAATAA AGATAAAATGG TACAATGATT	3600
TGCGTCGTTA CATCGAAATT ATCATGCAGA CAGACATGAT TCGTGAGAAA AAATAAAAG	3660
TGACTAACGA AATCACGAAT GCTATGGAAT ATTATAACAG CTCCTTTTG AAAGCTGTAC	3720

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CTCATTGAC GACGGAGTAT AAGCGCTTAG CGCAAGCGCA TGGTCTGAAT TTAAAACAGG	3780
CTAAACCAAT CACCATGGGT ATGTGGATAG GTGGTGACCG TGATGGAAAT CCATTTGTTA	3840
CAGCAAAGAC CTTGAAGCAG TCTGCACTCA CTCAGTGTGA AGTCATCATG AACTACTATG	3900
ATAAAAAGAT TTACCAACTT TATCGTAAT TTTCTCTTTC AACTAGCATT GTCAACGTCA	3960
GCAAGCAAGT CAGAGAAATG GCTCGTCAAT CCAAGGATAA CTCGATTTAC CGCGAAAAAG	4020
AGCTTTACCG TCGTGCCTTG TTTGATATTCA AATCAAAAAT TCAGGCAACT AAAACCTATC	4080
TGATTGAGGA TGAAGAAGTT GGGACTCGTT ATGAAACCGC CAATGATTTC TACAAGGATT	4140
TGATTGCCAT TCGAGATTCT CTACTAGAAA ATAAGGGCGA GTCCTTGATT TCAGGTGATT	4200
TTGTGGAATT ATTGCAGGCA GTAGAGATAT TTGGTTTTA CTTAGCATCA ATTGATATGC	4260
GACAAGACTC TAGCGTCTAT GAAGCCTGTG TGGCAGAACT CTTGAAATCA GCAGGAATT	4320
ATTCTCGTTA TAGCGAGTTG AGCGAAGAAG AAAAGTGTGA CCTTCTCTTG AAAGAATTAG	4380
AAGAAGATCC CCGAATTCTT TCTGCGACTC ACGCAGAAAA ATCAGAATTAA TTAGCAAAAG	4440
AATTAGCTAT TTTTAAGACG GCTCGTGTG TGAAAGATAA GTTGGGAGAT GATGTCATCC	4500
GTCAGACCCT CATTTCACAT GCAACCAGCC TTTCTGATAT GCTAGAATTAA GCTATTCTGT	4560
AAAAAGAAGT AGGACTGGTG GATACGGAAA GGGCGCGTGT TCAGATTGTT CCCCTTTTG	4620
AAACAATTGA AGACTTGGAT CATTCAAGGG AAACAATGAG AAAATATCTT TCTCTTAGCC	4680
TTGCCAAAAA ATGGATTGAC TCACGAAATA ACTACCAAGA AATCATGCTT GGCTACTCTG	4740
ACAGTAATAA AGATGGCGGT TACTTGTCA CATGTTGGAC CCTCTACAAG GCTCAACAAC	4800
AATTGACTGC TATTGGAGAT GAATTGGCG TTAAGGTTAC CTTCTTCCAT GGCGTGGTG	4860
GTACTGTCGG TCGTGGTGGT GGGCAACCT ATGAAGCCAT TACATCTCAA CCGCTCAAGT	4920
CTATCAAGGA TCGTATCCGC TTGACGGAGC AGGGTGAAGT AATTGGGAAT AAATACGGTA	4980
ACAAAGACGC CGCTTACTAT AACCTGAAA TGCTAGTATC GGCAGCTATT AACCGTATGA	5040
TTACTCAGAA GAAGAGCGAT ACCAATACCC CAAATCGTTA TGAAACCATT ATGGATCAAG	5100
TAGTGGACCG TAGTTACGAT ATCTACCGTG ATTTGGTCTT TGGTAATGAG CATTCTATG	5160
ATTATTTCTT CGAGTCAGT CCAATCAAGG CTATTTCAAG TTTTAATATT GGTTCTCGTC	5220
CAGCCGCTCG TAAGACTATT ACTGAAATCG GTGGTTTGC G TGCCATCCCT TGGGTATTCT	5280
CATGGTCACA GAGTCGTGTT ATGTTCCCTG GATGGTACGG GGTTGGTTCA AGCTTCAGG	5340
AATTATCAA TAAAATCCA GAGAATATTG CTATCTTACG AGATATGTAC CAAAATTGGC	5400
CTTTCTTCCA ATCGCTTCTT TCAAATGTTG ATATGGTTT GTCAAAATCA AATATGAATA	5460
TTGCTTTGA ATATGCTAAA CTTTGTGAAG ACGAGCAAGT TAAGGCCATC TATGAGACTA	5520

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TTTTAAATGA ATGGCAAGTT ACTAAGAACG TTATCTTGGC TATTGAAGGA CATGACGAAC	5580
TCTTAGCTGA CAATCCATAT CTAAAAGCTA GTCTGGATTA CCGTATGCCT TACTTTAATA	5640
TTCTCAACTA TATTCAGTTG GAGTTGATTA AACGCCAACG TCGTGGAGAA TTGTCCAGTG	5700
ATCAAGAACG ATTGATTCAT ATCACCATCA ACGGAATTGC GACAGGATTG CGTAATTCA	5760
GTTGATAATT TTCAAGAGTG AATGCTAAAA GTGAATATCA AAAAAATTCT AATAGACTAT	5820
TGACAAGTAG TTTAAAAATG ATATAATTCA ACCATTCAAGA AAAGTAATCA TACAAACTTT	5880
TTAGAGAGTC TGTGGTAGCT GAAAACAGAT AAGTGGCAAT GATGAAAATT GGGCTGAATG	5940
CTATTTAGAA TTTGAAATTA TAAAAATTG GTAAGCACAC CTTACAGTGC ATCTCGTTAT	6000
TGCGAGACTG AGCGATAGGG AAATTCCCTA TAATTGAGGT GGTACCGCGC ATCGACGTCC	6060
TCACACAAAGT TTTTGTGTG AGGATTTTT TGATGGAGGT TAGTATGGAA AGAAAACGAT	6120
GGCGTCGCTT GTTTAGATAA GTGAAATATG TAAAGGAAA TAAAAAGGAG AAACAGAATG	6180
AAAAATAAAC GTTTAATTGG AATTATTGCT CCATTAGCAG TCTTAGTGC AGGAAGCTTG	6240
ATTTATTCTT CAATGAATAA ATCAGAAGCT CAGAATAATA AGGATGAGAA GAAAATAACC	6300
AAGATTGGTG TGCTTCAATT TGTGAGCCAT CCATCCCTTG ATTTGATTAA TAAAGGGATC	6360
CAAGATGGAC TTGCAGAAGA AGGATATAAA GATGATCAAG TTTAAATTGA TTTTATGAAC	6420
TCAGAAGGTG ACCAAAGTAA GGTTGCGACA ATGAGTAAAC AATTGGTTGC AAATGGAAT	6480
GACCTTGTGG TTGGTATCGC AACACCAGCA GCCCAAGGGT TGGCTAGTGC AACAAAAGAC	6540
CTACCGGTTA TCATGGCCGC TATTACAGAC CCAATTGGTG CTAACCTGGT TAAAGATTG	6600
AAAAAACCAG GTGCCAACGT TACAGGGGTA TCTGACCACA ATCCAGCTCA ACAACAAGTT	6660
GAACTCATCA AGGCTCTGAC ACCGAATGTG AAAACAATCG GAGCTCTTTA CTCAAGTAGC	6720
GAAGACAATT CAAAAA	6735

(2) INFORMATION FOR SEQ ID NO: 105:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6516 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 105:

CTAGAGGATC CCAGCAGGTA AATTGGCTTC AGCTGGCAAA AAAGTTGCC TCGTTGAACG	60
CAGCAAGGCT ATGTACGGTG GAACTTGTAT CAACATTGGT TGTATCCAA CTAAAACCTT	120

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GCTAGTTGCT	GCTGAAAAGG	ACTTGTCTTT	TGAAGAAGTC	ATTGCTACTA	AAAACACGAT	180
CACTGGTCGC	CTCAACCGTA	AAAACATATGC	GACTGTTGCT	GGTACAGGCG	TAGATATCTT	240
TGATGCGGAA	GCTCACTTCC	TTTCAAATAA	AGTCATCGAA	ATCCAAGCTG	GTGATGAAAA	300
GAAAGAACTG	ACTGCTGAA	CAATCGTCAT	CAACACTGGT	GCTGTTCAA	ACGTCTGCC	360
AATCCCTGGA	CTTGCTACAA	GCAAAAACAT	CTTGACTCA	ACAGGTATCC	AAAGCTTGGA	420
CAAATTACCT	GAAAAACTTG	GAATCCTTGG	TGGCGGAAAT	ATCGGTCTG	AATTGCCGG	480
CCTTTACAAC	AAACTTGGAA	GCAAGGTAC	AGTCCTAGAT	GCCTTGGATA	CATTCCCTACC	540
TCGTGCAGAA	CCTTCCATCG	CAGCTCTTGC	TAAAACAATAC	ATGGAAGAAG	ATGGCATTGA	600
ATTGCTTCAA	AATATCCATA	CTACTGAAAT	CAAAACGAT	GGTGACCAAG	TGCTTGTGCGT	660
AACTGAAGAC	GAAACTTACC	GTTCGACGC	CCTTCTCTAC	GCAACTGGAC	GCAAACCAAA	720
TGTAGAACCA	CTTCAACTTG	AAAATACAGA	TATTGAACTA	ACTGAACGTG	GTGCTATTAA	780
AGTAGACAAA	CACTGTCAAA	CAAACGTTCC	TGGTGTCTTT	GCAGTTGGAG	ATGTCAACGG	840
TGGCCTTCAA	TTTACTTACA	TTTCACTTGA	TGACTTCCGT	GTTGTTACA	GCTACCTTGC	900
TGGAGATGCC	AGCTATACAC	TTGAAGACCG	TCTCAATGTG	CCAAATACTA	TGTTCATCAC	960
ACCTGCACCT	TCACAAGTTG	GTTCGACTGA	AAGCCAAGCA	GCTGATTGTA	AACTTCCATA	1020
CGCTGTTAAG	GAAATCCCCG	TTGCAGCAAT	GCCTCGTGGT	CACGTAAATG	GAGACCTTCG	1080
CGGTGCCTTC	AAAGCTGTTG	TCAATACTGA	AACAAAAGAA	ATTCTTGGAG	CAAGCATCTT	1140
CTCAGAAGGT	TCTCAAGAAA	TCATCAACAT	CATCACTGTT	GCTATGGACA	ACAAGATTCC	1200
TTACACTTAC	TTCACAAAAC	AAATCTTCAC	TCACCCAACC	TTGGCTGAGA	ACTTGAATGA	1260
CTTGTGCGG	ATTAAAGTTG	AGATTTAAC	GTATCGAAC	GCCCTCTTG	GGCTGTTTT	1320
ACTTCTGCGG	AATCTCAAAT	CTGTCTTCT	CCTCTTTAT	GATATAATAG	AAACATGAAC	1380
TTAAAAACTA	CTTTGGGCCT	TCTTGCTGGG	CGTTCTTCCC	ACTTCGTTTT	AAGCCGTCTT	1440
GGACGTGGAA	GTACGCTCCC	AGGGAAAGTC	GCCCTTCAAT	TTGATAAAGA	TATTTACAA	1500
AACCTAGCTA	AGAACTACG	GATTGTCGTT	GTCACTGGAA	CAAATGGAAA	AACCCTGACA	1560
ACTGCCCTCA	CTGTCGGCAT	TTTAAAAGAG	GTTCGAAACA	ACCTTCCTAA	CAGCCAAATC	1620
GGTGCCAAACA	TGATTACAGG	GATTGCAACA	ACCTTCCTAA	CAGCCAAATC	TTCTAAAAC	1680
GGGAAAATA	TTGCCGTCT	CGAAATTGAC	GAAGCCAGTC	TATCTCGTAT	CTGTGACTAT	1740
ATCCAGCCTA	GTCTTTTGT	CATTACTAAT	ATCTTCCGTG	ACCAAGATGGA	CCGTTTCGGT	1800
GAAATCTATA	CTACCTATAA	CATGATATTG	GATGCCATTC	GGAAAGTTCC	AACTGCTACT	1860
GTTCTCCTTA	ACGGAGACAG	TCCACTTTTC	TACAAGCCAA	CTATTCCAAA	CCCTATAGAG	1920

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TATTTTGGTT TTGACTTGGA AAAGGGACCA GCCCAACTGG CTCACTACAA TACCGAAGGG	1980
ATTCTCTGTC CTGACTGCCA AGGCATCCTC AAATATGAGC ATAATACCTA TGCAAACCTG	2040
GGTGCCTATA TCTGTGAAGG TTGTGGATGT AAACGTCTG ATCTCGACTA TCGTTGACA	2100
AAACTGGTTG AGTTGACCAA CAATCGCTCT CGCTTTGTCA TAGACGGCCA AGAATACGGT	2160
ATCCAAATCG GCGGGCTCTA TAATATCTAT AACGCCCTAG CTGCTGTGGC CATGCCCGT	2220
TTCCTAGGTG CCGATTCGCA ACTCATAAA CAGGGATTTG ACAAGAGCCG TGCTGTCTT	2280
GGACGCCAAG AAACCTTCA TATCGGTGAC AAGGAATGTA CCCTTGTCTT GATTAAAAAT	2340
CCAGTCGGTG CAACCCAAGC TATCGAAATG ATCAAACTAG CACCTTATCC ATTTAGCCTA	2400
TCTGTCTCC TTAATGCCAA CTATGCAGAT GGAATTGACA CTAGCTGGAT CTGGGATGCA	2460
GACTTTGAAC AAATCACTGA CATGGACATT CCTGAAATCA ACGCTGGCGG TGTCGTCAT	2520
TCTGAAATCG CTCGTCGCT CCGAGTGACT GGCTATCCAG CTGAGAAAAT CACTGAAACG	2580
AGTAATCTGG AGCAAGTTCT CAAGACCATT GAGAATCAAG ACTGCAAGCA TGCCTATATT	2640
CTGGCAACTT ATACTGCCAT GCTGGAATTT CGTGAACACTGC TGGCTAGTCG TCAGATTGTT	2700
AGAAAGGAGA TGAACTAATG GTTTATACTT CACTTCCTC AAAAGATGGC AATTACCCCT	2760
ATCAGCTCAA CATTGCCAC CTCTACGGAA ATCTCATGAA TACTACGGGG ACAATGGAAA	2820
CATCCTCATG CTCAAGTATG TGGCTGAAA ACTGGGAGCC CATGTGACCG TTGACATCGT	2880
TTCTCTCCAT GATGACTTTG ATGAAAATCA CTACGACATC GCCTTTTCG GTGGTGGTCA	2940
AGACTTTGAA CAAAGTATCA TTGCAGACGA CCTACCTGCT AAAAAAGAGA GCATTGACAA	3000
CTACATCCAA AACGACGGTG TAGTTCTGGC TATCTCGGGT GGTTCCAAC TATTGGTCA	3060
ATATTATGTT GAAGCTTCAG GAAAACGTAT CGAAGGGCTA GGGGTCATGG GACACTACAC	3120
GCTCAACCAG ACCAATAACC GTTTTATCGG TGACATCAAG ATTACAAATG AAGATTTCGA	3180
TGAAACCTAC TATGGATTTG AAAATCACCA AGGTCGTACC TTCCTCTCTG ATGACCAAAA	3240
ACCGCTGGGA CAGGTTGTCT ATGGAATGG AAACAAACGAA GAAAAGGTG GTGAAGGGGT	3300
TCATTATAAG AATGTCTTTG GTTCTACTT CCACGGGCCT ATCCTCTCTC GTAATGCCAA	3360
TCTGGCTTAT CGCCTAGTTA CTACTGCCCT CAAGAAGAAA TATGGTCAGG ACATCCAAC	3420
CCCTGCCTAT GAGGACATTC TCAGCCAAGA AATCGCTGAA GAGTACAGTG ACGTCAAAAG	3480
CAAGGCTGAC TTTTCTTAAA CAAAGGAAAA TGATATCAA GAACTCCGTT ATCTGTGCG	3540
AGTTTTTGT CTTTCTTT ACCCTCTCC CTTGCATTT CTCTCATTTC TTGCCAAAT	3600
AGAGGGGTAG AAAGAAGGTA GCATATGTCT AAATTACAAC AAATCCTAAC ATATCTGAA	3660

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TCAGAAAAAC TAGACGTCGC TGCGTATCT GACCCCGTCA CAATCAATTAA	CCTCACTGGT	3720
TTTTACAGTG ATCCCCATGA ACGCCAAATG TTCCCTCTTG TCCTAGCAGA TCAGGAACCT		3780
CTCCTCTTG TCCCAGCTCT TGAAGTAGAA CGTGCAAGTA GCACCGTTTC CTTCCCAGTA		3840
GTGGGCTATG TCGATTCTGA AAATCCATGG CAAAAAAATCA AACATGCTCT TCCACAACCTT		3900
GACTTCAAAC GTGTCGCTGT TGAGTTTGAC AATCTCATCT TGACCAAATA CCATGGTTTG		3960
AAAACAGTTT TTGAGACTGC TGAGTTTGAC AACCTCACTC CTCGTATCCA ACGCATGCGC		4020
CTCATCAAAT CAGCTGATGA AGTGCAAAAAA ATGATGGTTG CAGGTCTTTA TGCTGACAAG		4080
GCTGTTCATG TTGGTTTGAA CAATATTCT CTTGATAAGA CTGAGACAGA TATCATCGCA		4140
CAAATCGACT TTGCCATGAA ACGTGAAGGT TATGAAATGA GCTTGATAC CATGGCTTG		4200
ACTGGTGATA ATGCTGCGAA TCCACACGGC ATTCCAGCAG CTAATAAGGT TGAAAATGAT		4260
GCTCTTCTCC TCTTGACCT GGGTGTCTG GTCAATGGCT ATGCGTCAGA TATGACTCGT		4320
ACAGTCGCTG TCGGCAAACC AGACCAAATTC AAGAAAGATA TTTACAACCTT GACTCTTGAA		4380
GCCCAACAAG CTGCTCTGAA CTTTATCAAG CCAGGTGTGA CTGCTCATGA AGTGGACCGC		4440
GCTGCCGTG AGGTCACTCGA AAAAGCTGGT TATGGTGAGT ACTTCACCCA CCGTCTCGGG		4500
CATGGTATCG GTATGGATGT CCATGAATTC CCATCTATCA TGGAAGGAAA CGACATGGTC		4560
ATCGAAGAAG GCATGTGCTT CTCTGTTGAA CCAGGTATCT ATATCCCTGG TAAAGTCGGT		4620
GTTCGTATTG AAGACTGCGG TGTTGTTACC AAGGATGGCT TCAACCTCTT TACAAGCACC		4680
AGCAAAGATT TGCTTTATTT TGATTAACCT ATATAGCCCC TATGCTTCC TTTCAAAATA		4740
TCTAGGGCT ATTTTATTGT CATTCTCTG CTATTATGCT AAAGAAATTG GCTGCAATAA		4800
TCTAACCTCA AGTGTCTGGA ATGATAACGA GGGTGTCTC CGCTTTATC AAAGACAAGG		4860
GATGAAACCC CAAGAAACAA CAATGGAAAT GATAATTGAT TAAGAAGTCA TCTATCAAAA		4920
GATGTTAGAA AAAGTTCAAT TTCACTAGAA AATGAGGAAA ATCTCCCCAC AATAAACACG		4980
ATAGTATCAG GTATTGTGTA CTGACCCCAA ACAGTTAGAC AATTAATTAA TCCGAAGGAT		5040
TAGTTCTGT ACTGCACAGG ACTAAGTCCT TTTAGTTTA CCTTAATTG TTTGTTGTTG		5100
TAGTAATCAA TATAGTCTAT AATGACTTGT TCCAATTGGT TAAGTGATT AAATGTTTC		5160
TCATAGCCAT AAAACATTTC GGATTTAAA ATGCCAAAGA AAGATTCCAT CATAACGTTG		5220
TCTGGCTGT TTCCCTGCG TGACATAGAT GCTTGAATTG CCTTATTCTC TAGGAACCGA		5280
TGATAAGAAT CGTGTGGTA TTGCCAGCCT TGCTCACTAT GGAGAATCGT ATTCTCGTAG		5340
TGCTTCTCTT TGAATGCCTG TTCCAACATT GTTTGTACTT ATTCTAAATT AGGCGAACAA		5400
GAAAGATTAA AAGCAATAAT TTCGCTGTTA AAGCCATCTA AAACCTGGTGA TAAGTAAAGC		5460

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TTTTGAGTAC TTGCTGGAAT GGCAAATTCA GTCACATCTG TGTAGCACTT TTCCATTGTT	5520
TTAGAGCCTT CAAATTGGGC TTGAATGAGA TTCTCTGCCT TCTTACCAAC GTCTCCTTTA	5580
TGAGAAGAAC ATTTCGTTT CTTCGCATT TTAGCTGTA AATTGAGTAC TTTCATCAAG	5640
CCTTGAACTC TTTTATGATT TACCAGATAA CCACGATTTC TTAGTTCTAA ATGAACCCGG	5700
CGATAAGCAT AATTCCCTT GTGTTCGATA AAGATGGATT GAATTTCACT TTTAAGCTCT	5760
TGGTCTTAT CTGTTTGTC TAGCTGTTTC AAGTGATAGT AGTAGGTCCA ACGAGCTAGT	5820
TTAATGGCTT CTAGAAGAAG ATCTAACGAA AACTCAGTCA TTAATTCTTG AACAAATTCT	5880
GTCTTCTTC TTTCTCTTT TCCTCCTTCA ATCGGAGTTC TCTTAACCTT TTTAGGATGG	5940
CATTCTCCGC TCTCAGGTAC TCTCCCTCTT GTTTCTCAA CAATAGTATA CCCGTTTTC	6000
CTGTATTGTG CTAGCCAGTT AAGAAGTATC GTACGACTTG GGAGACCGTA TTCAAGAGAA	6060
ACTCTATCTT TAGTCCAGCC TTCATGTCAG ACTTTATTAA CCCCAATTAT TCACCCAAA	6120
TCTAAAAACC ATCCAGAAC CTTGCCCTAG CTTAGATCCT GGATGGTTTC TTTTTTCACC	6180
CAATGGGTGT TTTTACTAG ACAAAAAAGA GTTCCCCTT TATGGTATAA GTGTAGAAAA	6240
AAACACAAAA AGAAAGGAAA CTCACATGAA CAGTTTACCA AATCATCACT TCCAAACAA	6300
GTCTTTTAC CAACTATCTT TCGATGGAGG TCATTTAAC CAGTATGGTG GTCTTATCTT	6360
TTTCAGGAA CTTTTTCCC AGTTGAAACT AAAAGAGCGG ATTTCTAAGT ATTTAGTAAC	6420
GAATGAmCAA CGCCGCTACT GTCGTTATTC GGATTTCAGAT ATCCwtGTCC AGTTCCCTTT	6480
TCAACTGTTA ACAGGTTATG GAACGGAATA TGCTTG	6516

(2) INFORMATION FOR SEQ ID NO: 106:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14654 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 106:

TTTTCAACCC ATATCGTGGC TCCTGAATAC TACTTACTGA CAACTATGCT ATCAGAGACT	60
TCTCTACTTG TTTTCTATAT CATTTCATC CATAGAAAAC AACTCATCCA CTTGGGACAT	120
ATCTTTAGCT ATACTGTTCG ATACTCTCTC TTTTCACTTT CCTTTGTAGC AATTTATTTC	180
CTGATTAATT TCGTGTATCC TGTAGATATG GTCATTAATT TGCCATTTT GATTAATACT	240
GGTTTGATTG TCTTGCTATC AGCTATCTCT TATATTAGTC TACTTGTCTT CACAAAAGAT	300

782	
AGCATTCT ATGAATTTT AAACCATGTC CTAGCCTAA AAAATAAATT TAAAAAATCA	360
TAGGAGTTA AAATGAAACA ACTAACCGTT GAAGATGCCA AACAAATTGA ATTAGAAATT	420
TTGGATTATA TTGATACTCT CTGTAAAAG CACAATATCA ACTATATTAT TAACTACGGT	480
ACTCTGATG GGGCGGTTCG ACATGAGGGC TTTATCCCTT GGGACGACGA TATTGATCTG	540
TCCATGCCA GAGAAGACTA CCAACGATT ATTAAACATT TTCAAAAGGA AAAAAGCAAG	600
TATAAGCTCC TATCCTTAGA AACTGATAAG AACTACTTTA ACAACTTTAT CAAGATAACC	660
GACAGTACGA CTAAAATTAT TGATACTCGA AATACAAAAA CCTATGAGTC TGGTATCTT	720
ATCGATATTT TCCCTATAGA TCGCTTGAT GATCCTAAGG TCATTGATAC TTGTTATAAA	780
CTGGAAAGCT TCAAACGCT GTCTTCAGT AAACATAAAA ATATTGTCTA TAAGGATAGC	840
CTTTTAAAG ATTGGATACG AACAGCCTTC TGGTTACTCC TTGACCGGT TTCTCCTCGT	900
TATTTTGCAA ATAAAATCGA GAAAGAAATT CAAAATATA GTCGTAAAAA TGGGCAATAT	960
ATGGCTTTA TCCCTCAAA ATTTAAGGAA AAGGAAGTCT TCCCAAGTGG TACCTTTGAT	1020
AAAACAATCG ATTTACCCCT TGAGAATTAA AGCCTCCCTG CACCTGAAAA ATTTGATACT	1080
ATTTGACAC AATTTATGG AGATTATATG ACCCTACCAC CAGAAGAAAA ACGCTTCTAC	1140
AGTCATGAAT TTCACCGCTTA TAAATTGGAG GATTAGGATG CAATATTAG AAAAAGAAGA	1200
AATTAAAGAA ATTCAACTAG CCCTGCTGGA CTATATTGAT GAGACTTGTA AGAAACATGA	1260
TATTCCTTAT TTTCTCAGTT ATGGAACCAT GCTTGGAGCC ATCCGCCACA AAGGTATGAT	1320
TCCTTGGGAT GATGATATTG ATATTTCCCT TTATCGTGAG GATTATGAGC GTTTACTGAA	1380
GATTATTGAA GAAGAAAATC ACCCTCGCTA CAAGGTTCTT TCCTACGATA CATCTTCTTG	1440
GTACTTCCAT AATTCGCAT CGATTTGGAA CACTTCTACT GTTATAGAAG ACCATGTTAA	1500
GTACAAGCGT CATGATACCA GCCTTTCAT CGATGTCTTC CCAATTGATC GATTACAGA	1560
CTTGAGCATT GTCGACAAGA GCTATAAGTA TGTGGCTCTT CGTCAACTAG CTTATATCAA	1620
AAAATCACGA GCAGTTCACG GTGATAGCAA ACTAAAAGAT TTTCTTAGAT TATGTAGCTG	1680
GTACGCTCTC CGATTTGTCA ATCCCTCGCTA CTTTTACAAG AAAATTGATC AACTAGTCAA	1740
AAATGCTGTA ACCAACACTC CTCAATATGA AGGAGGAGTT GGGATCGGTA AGGAAGGGAT	1800
GAAAGAAATC TTCCCAGTTG ATACCTTTAA AGAACTGATT TTAACTGAGT TTGAGGGCCG	1860
TATGTTGCCT GTTCCCAAAA AATATGACCA ATTTTAACC CAGATGTATG GCGATTATAT	1920
GACACCACCA TCAAAAGAAA TGCAAGAGTG GTATAGTCAT AGCATTAAAG CTTATCGCAA	1980
AAACTGATTG AGGGGGATTA TACAAACTAC TAAGATAGAG GTTATTCAA AACATAATT	2040
TAGTAGAAAA TGAAATACAT ATTCCCCACAA TAAAACGCAT CATATCAAGG TTTTGAAAAA	2100

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ACCTTGATAAT GATGCGTTT ATAATTTAA AGACTTTTT CTATAGTAGA TTGAAATAAG	2160
ATGCGAACAA ATCAATTAGA AAATTCAAAT TAATTTATAG AAATATTTA GTATTCCTGT	2220
GTACTGTTCT AAATTCAAGTC TGCTATATCT TATTTTCTA TTTAAATCGC TTCTGTAACA	2280
AAGCTACGAC TTTCAAGTAC CTTAACCATG GCATTAGCTG TATCTAGCGC TGTGAAGAGG	2340
GGCACCCCGT GTTCAATGGC TGAACGACGA ATTTGCTCAC CATCTCGTC AGCAGTCGT	2400
TTTGTTCCTA CTGTGTTAAT GATAGCTTGA ATTCTTCCTT TGCACAAA ACTTGGGATA	2460
TCCTTATCGT CATCACCAAT CTTACCAACA GGTTGGGCTT GCAAGCCATG ACTAGCAAAG	2520
AAGGCTGCTG TCCCCTCTGT CGCAAGGATT CCATAACAA TGTTTGAA ACGACGAGCC	2580
AAGTTCAAGG CTTCTCTTT GGCATCATCA GCGATGGTAA AGACGACATT ACCAAAAGTT	2640
GGCAAGTGTGATAAGAAGC TTCAAAGGCT TTATAGAGAG CTTTTCCAAGTACATCA	2700
GAACCCATAA CTTCACCTGT TGACTTCATT TCAGGACCGA GCAAGCTGTC TACCTTAGCT	2760
AGTTTGGTAA AGGAGAAGAC AGGTGCCTTG ATATGAACAC GGGTGCTTC AGGGTAAAGT	2820
CCATTGGT AGCCAAGTTC TGATAAACTT TGACCAAGAA TGAGTTGGT CGCTACTTGA	2880
GCCATAGGAA TATTGGTTAC CTTAGATAGG AATGGAACAG TACGGCTGGC ACGTGGATTG	2940
ACCTCAATAA CGTAGACTTT TTCATCCTTG ATAACAAACT GGATGTTCAT CATTCCAAGG	3000
CAGTGAAGAC CGATTGCTAA GCGTTGGTG TAGTCTGGCA TGGTCTCCTG AACCTTTGC	3060
GACAAGGTTT GTGGTGGGTA AACAGCCATT GAGTCACCTG AGTGGACACC AGCACGTTCG	3120
ATATGCTCCA TGATACCAGG AATGAGTACA TTTTACCAT CTGAAATGGC ATCAACTTCG	3180
CACTCTGCC CAACGATATA AGAGTCGACA AGAACTGGGT GGTCTGGACT AGCCTTAACA	3240
GCAGTTCGCA TGTAAGAACG AAGGTCTTCT TCGTTTCAA CGATTTCCAT GGCACGTTCA	3300
CCAAGTACAT AAGATGGCG GACAAGAACT GGGAGCCAA TCTTGCAGC TGCAAGAGCT	3360
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GCTTGCTCGA AGAGGTACG GTCTCGGCA CGATCTAGGT CAGCAACCTG TGTACCAAGG	3480
ATGGTCACAC CTGCTTTGC CAATGGCTCC GCAAGGTTGA TGGCTGTTG ACCACCGAAC	3540
TGAACGATAA CTCCCTTGG TTGTTCCAAG TCAATGACGT TCATAACATC TTPGAATGTC	3600
AATGGCTCAA AGTAAAGCTT ATCTGATACA GAGAAGTCTG TTGAAACGGT CTCTGGGTTT	3660
GAGTTCATGA TGATAGCTTC ATAACCAGCT GCCTGGATAG CCTTAACAGA GTGAACGGTT	3720
GCGTAGTCAA ACTCAACCCCC TTGACCGATA CGGATTGGAC CTGAACCTAG GACAAGTACA	3780
GATTCTTAT CAGATCTGAT AGATTCAATT TCCCAACCAT AGGTTGAATA GAAATATGGC	3840

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GTTTCGGAGT CGAACTCTGC CGCACAAAGTG TCTACCACATCT TATAAAACTGG AACAAATCTTG	3900
TTTTCCAAGC GAAGTTGGCG AACTTTATCA TCAGTCGTTCC CCCAGAGTTCA AGCAATCTTA	3960
CGGTCTGAAA AACCATTAAAG TTTGGCTGTT TTCAAAACTT CTAAATCTTG TGGATGAGCA	4020
CCCAATTCTT GCTCAATTTC AAAGATATGC AAGAGTTTAT CAAGATAGAA GATATCAATT	4080
TTTGTAAAGCT CTGCAATTTC TTCAGGTGTG TAGCCACGAC GAATGGCTTC TGATACGTAG	4140
AAGAGACGGT CATCTTGGGC TTTGACAACC TTTCAATCA AGGCATCATC AGAAACTGCT	4200
GCAAGTTCAAG GTATTTCATT GTGGTGCACC CCAATTCAA GGGAGCGGCA GGCCTTGAGA	4260
AGAGATTCCCT CGATGTTACG ACCGATTGCC ATGACTTCTC CAGTCGCCTT CATTGTGTA	4320
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ACGTAGTCAA GGGCTGGTCA AAACATGGCA TAGGTTGAAC CTGTAACCTGG GTTTATAACC	4440
TCATCCAAGG TCAAACCTAC TGCAATCTTG GCAGCCAAC TAGCAATCGG ATATCCTGTC	4500
GCTTTAGAAG CAAGGGCTGA CGAACGTGAT ACACGAGGGT TTACTTCGAT AACATAATAC	4560
TTGAAGCTGT TAGGATCAAG AGCTAGCTGA ACATTACATC CACCTCAAT CTTGAGGGCA	4620
CGAATAATGC TCAAGCTCGC ATCACGAAGC ATTGGTTTT CATACTCTGA CATGGTTGC	4680
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GTTGCAGGAT TTGAGTTAAC CAAAACAACC TCATAACCTT CCTCTTTCAA CGACAAGCAA	5340
GCCTGAGTCC CAGCGTAGTC AAACTCAGCA GCCTGACCAA TAATAATCGG ACCAGAACCA	5400
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TTTTTAGCAC CGTCCGTAGC CCGTATTCAAG TTCAGCAAAT ACGGAGCACC CTTCTCCTT	5580
CTATTCTCGTCG CCTCTCAGGG CGACATTAAA TAAGATACAA AGGACGAATA GAAAGCGATT	5640

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GAATTTAGG AAATCAAGGA AGGATTGACA ATCCAAGTTG GTTTCTCTAC ATTCTGAGCT	5700
TTCCGTCCGT GTTCAGTTAC ATAAATTCTC CGACGAGCTT TTACTCGTTC TTAGTTGAT	5760
TGTTTAAAAA CTTCCATCAT CTCGATAAAC TCGTCAAATA GGTAGCTAGC GTCGTGTGGC	5820
CCAGGAGCTG CATCTGGGTG GTATTGAACA GAGAAAGCAG GTTGGTATCT GTGGCGCACA	5880
CCTTCCACTG ACTTGTCAATT GATTCTTCG TGCGTAATAA TCAAGTGCTC TGGCAAATCC	5940
TCGCGGCTGA CTGCATAACC ATGGTTCTGG CTGGTGAAGT CTACTCGTCC TTGCGCGATT	6000
TCACGTACCG CATGGTTGAA TCCACGGTGG CCAAACCTCA TCTTATAGGT CTTAGCCCCG	6060
TTTGCCATTG CAAAGAGTTG GTGTCCCATA CAAATACCAA AGATTGGAAT TTTCCTTGT	6120
ACACCGCGAA TCATGTCGAG TGCTTGTGGA ACGTCTTCTG GGTTACCTGG ACCATTGAC	6180
AACATAACTC CGTCAGGATT GAGATGGAGA ATTTCTTCAG CCGTTGTCCG ATAAGGAACA	6240
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TCCACTAGCA CCACGCTCAA ACCAACCTCT GGAGCTGGAT AAGACGTTT AGTAGAAACC	6360
TGTTTGATAT TGTCTGTCGG TAAAACTGTT GCTTGGAGCT GGTCCGTAC ATGGTCCATA	6420
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CCGGTCATCC CTGTATTTAA GACGATTTCG CCTGTTACAT CAATATCTGC TCCGAAGGCC	6780
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ACGGAGAAC ATCATGACAT CAACCTGATC AATGATTTCA TCAATGGTTA CAAACTGTCC	7200
ATAGTCTGCA AACTCTTGAC TTCTCCATTG CTCAGGTCCA GCGAAAAAGA GTTCAGCTCC	7260
CAAGCGTTTC AAAATCTGCA TATTGGATTT GGCAACGCGT GAGTGGTCCA AGTCACCTGC	7320
AATAGCAACT TTAAGACCCCT CAAAGTGGCC AAATTCCCTCA TAAATGGTCA TCAAATCAAG	7380

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CAAGCTCTGG	CTAGGGTGT	GGCCCGAAC	ATCTCCACCA	TTGATGATGG	AAGTCGTAAT	7440
CGTTGGACTA	GCAATCAATT	CTCTATAGTA	GTCGACCTCT	GGATGGCGAA	TCACACAGAC	7500
ATCCACTCCT	AAAGCAGACA	GAGTCAAAAT	GGTGTCAAA	AGTGTCTCAC	CCTTATTAAC	7560
CGAGCTAGTC	TTCACATCAA	AGTCAAGTCG	TTCCAATCCA	AGTTTAATCT	CTGCGACTTC	7620
AAAGGACTTA	TGTGTCCGTG	TAGAATCCTC	AAAGAAGAGA	TTGGAAACAA	TCGGATGGTC	7680
TTCATAGGGA	AGCTGGGCTC	CATTTTAAA	CTCAATTCC	CGCTTGATCA	ATTCATTAC	7740
TTGATCGACA	GTGAGGTCTT	CCATGGACAC	CACATGGTTC	AATGCTTGT	GATTTCTGA	7800
CATGGCTACT	CCTTTAACTT	TCTAAGCTTC	TTCAAGTATC	AGAACTCTGT	CTTGGTCATC	7860
AAAGTCTGTC	ATCTCTACGA	TGATTTCTTC	AGAACGACTG	GTTGGGATAT	TTTTCCAAC	7920
GTAATCTGGA	CGGATTGGCA	ATTCTCTATG	TCCACGATCG	ACTAGAACTG	CTAAACTCAC	7980
ACCGGCAGGA	CGACCATGAC	CGACAATATT	ATCAATAGCA	GCACGGATGG	TACGACCTGT	8040
ATAGAGCACA	TCATCCACCA	AGATAACTTC	GCGGTCTGTC	ACATCGACAG	AAACCAAAGA	8100
AGTATCTTCT	CCACTTTAA	CATCATCACG	GAAAGGTTA	GTATCCAATT	CCACAAACAGG	8160
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ACGAGTTTA	ATACCAGCCA	AGACGATCTT	ATTCAAATCT	TTGTTGCGTT	CGATAATCTC	8280
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CATGACAAAC	CTCCAAAAAG	AAAAGTCTCC	TTAAACAAGG	AGACTTGAAA	TTTATAGCCA	8400
AGCGAGCCCT	ACTGCACACA	GTATAGACTT	CACCTTCTA	CTTTATCGCG	CTCCTTGCCT	8460
GCCTCACGGG	ACAGGTTAA	AGGAATATT	AGTTATCATT	TACTATAGCA	CAAAGCATGC	8520
TTAAAATCAA	GCAAAAGT	TCAATGTAGC	ATCTTACAAA	TTGCTAAAAT	CATAATAATTG	8580
TGGGTACTGG	TCACACTCTG	GATTTTTGG	ATGGCAAATG	GCTCTTCCAA	AATAAATCAT	8640
GGCCTGATGG	GCAGCTAAC	ACTGCTCAGG	CGGCAAGATA	TCCATGACCC	GCTTTTCCAC	8700
CTCAAGTGGC	GTCGCTGATT	TTTGACAAT	ATCGTGGTGT	TTGCAAATAC	GCTCCACATG	8760
AGTATCCACT	GCAAAAGCTG	GAATTCCAAA	TCCTACACTC	ATGACAACAT	TGGCTGTCTT	8820
GCGACCAACA	CCTGCCAAC	TCTCCAATTC	TTCACGTGTC	TGAGGGACTT	GACCATCAA	8880
ATCGTCTAGT	AACTGTTGGG	CACATTTTT	AAGGAATTAA	GCTTTATTCC	GATACAATCC	8940
CAAGCGAGAA	ATATGTGAAG	CAATCTCACT	CTCTGTCGCT	ACAGACATAG	CTTGGGGTGT	9000
TGGAAAGGCA	ACAAAGAGAC	CTGGTGTGGC	CTTATTTACC	GCTGCATCTG	TCGTCTGGC	9060
TGATAAACATG	ACCGCAACCA	GGAGTTCAA	ATGATTGGTA	AAATCAAGAC	TAGGCTTGGC	9120
ATCTGGGAAG	AGGGCAATGA	TTTCTTCTAG	CACCTTCGT	GCTCGTTTT	TTGACAAGAC	9180

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CATTATTCAAT	CTCCGTCAAA	TAGTCCTTGT	AAGGCCAGCAA	AAGGACTGTT	TTCTTCTTTC	9240
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ATAAACCTT	GACCAGCTTC	TTCTTCAGCC	GTCAAGACCT	TGATAGGAAT	GTTCAGCAGG	9360
ATATTGTCTG	ATACACTCTC	AGCAAGGTCA	AGCTCCCCAT	TTTCGATGGG	CAAGACCAAG	9420
TCATCATCTA	AAACTTCTTG	ATCTAGCTGG	TTAGTTGC	CTTCCATGAA	AACTTCCGTG	9480
ACTGGATAAG	ATTCAACTAA	CTCAACTGGC	TCCACTACTGC	GACTCGACGC	AAGAACAAATG	9540
GTATAAGATA	GTTGATAATC	TAAGAAATAC	ATACGGTCTT	CATATTGTAC	TTTCCCAACT	9600
GCAAGGATAT	CTTTTACATC	TAAAATTCT	TGATTACGTG	CACGCAGGTC	ATCAACTAAA	9660
TCTAACGTTT	GTTCAAAGTT	CAAACCTTCA	GACTGCTTAC	GAATTTCTTG	AATATTTAAT	9720
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GCACACCAAA	CTTTGTAATT	AAAATTCAAA	ATTTTAACAT	ATTTACTATG	ATAGTTTTAT	9840
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GACAAGAAC	TCATTCAACT	CCTATCCAAG	TTAAATAAAA	GCTACCAAAA	CTGTAAACAG	9960
GGTACGGCAG	ATGATATTG	ACTACAAGAG	CTGCTAAACA	CTACTATGCA	AGAGCTCAA	10020
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AGTCTTCTGA	TTGGACTGGG	TAGCCTAAAA	CTAAACGATC	AAGCACGGAC	TGCTTGGCGA	10140
AACTATGATA	AATTCCATTA	CGATCATGTC	AAACACGTAC	TAAGTCTCTA	TGGACCTGTT	10200
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ATATAAGAT	ACTAATACTC	GGAGGTAAGG	GAGACATGAA	CAACTAAGTC	TATCAAATAA	10320
AGAACCTTTA	TTTAGTAGAT	CTTGTTTTG	TCTCTTTTG	TGTGCTCTT	TATGCTCTT	10380
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GCTTTTGTT	ATGCACTATG	AACATTCTAG	AAAGGGAAAT	CATATGATAA	AAATCAATCA	10500
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ACACGACTAC	TTCTTTTAG	ATTCTATTGA	TTTAGACTAC	AGTATCCTCT	ATCGTTGGC	10800
GGAGGAATTG	CATTTGATA	GCAATCGTT	CGCAAGTGAC	CAAGAGATTG	GCAATCTATC	10860
AGGGGGCGAA	GCTTTGAAAA	TTCAGCTTAT	CCATGAGTTA	GCCAAACCT	TTGAGATTCT	10920

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ATTTTTAGAT GAACCTTCAA ATGACCTAGA CCTTGAGACCA GTTGATTGGC TAAAAGGCCA	10980
GATTCAAAAG ACCAGGCAGA CCGTTATTTT CATTCCCCT GATGAAGACT TTCTTCTGA	11040
AACGGCAGAC ACTATTGTTC ACTTGCGACT GGTCAACAC CGTAAAGAAG CGGAAACGCT	11100
AGTAGAGCAT TTAGACTATG ATAGCTATAG TGAGCAGAGA AAGGCTAATT TTGCCAAACA	11160
AAGTCAGCAA GCTGCTAACCA ACCAAAGAGC CTACGATAAA ACCATGGAAA AACATCGGAG	11220
AGTTAAGCAA AATGTAGAAA CTGCGCTTCG AGCTACCAAA GATAGTACTG CCGGTCGCCT	11280
ATTGGCTAAA AAGATGAAAA CTGTCCTCTC ACAAGAAAAA CGCTACGAAA AGGCAGCTCA	11340
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ACCATTACCA GCTTCTAAAG TCTTAGTCCA ACTGGAAAAA GAAAATTGT CCATTGACGA	11460
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AGAAGTCTGC TCGATCATCT ATCGCATGAC AGAACACGGT TTGAAGCTAG TTAATTAGA	12000
AGATTTATAA ATTTGCAACA TAGAAAAAT CCAGAGACGA CCTCTGGATT CTTTTACATC	12060
TGTTTTAAAC GTTCAATCCG TTCTGAGATA GGTGGGTGGG TATAAAAGAG TTTTTGGAAC	12120
CCCCCACCTT TCTTAGGATC ATTGATATAA AGGGCACTGC TAGCATCATC GACGTGGCGA	12180
CTCATAGGTT TGCTATTGTC CAACTTATCT AGGGCATTAA TCATTCCCTG GGGATTGCGA	12240
GTCAGCTCGA CACTAGATGC ATCTGCCAGA AATTCCCTCT GACGAGAAAT AGCGAGCTGA	12300
ACCAAGGTG CAGCGAGAGG TGCCAGTACA ATAGCTAGTA GGGAAACAC TAGCATAATG	12360
ATTTCAAGAC CATTCCATC TCGGTCTCATCA TCACCTCGTC TGCGACCTGC TCCACCCAC	12420
CACATCATAC GACCTGCCAT ACTAGAAAGC ATGGTGATAG CACTAGCAAG GGCAACTGCA	12480
ATAGTCGAAA TACGGATATC ATAATTACGA ATATGACTGA CTTCATGTC CATAACAGCT	12540
TCTAGTTCTT CACGATTCT GATAGCTAGT AGACCTGAAG TCGCAGCAAC AGCCGCATTT	12600
TGAGGATTAG AACCTGTCGC AAAGGCATTT AAGGCTGGAT CATCAATGAT GAAAACACGG	12660
GGCATAGGAA TCTGAGCGAC CAGAGCCATA TCTTCCACTA CATGGTAGAG GTCTGGTGC	12720

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GTTTGCTCAT CCACCTCACG CGCTCCATTG ATGGACATGA CAATCTCTGT CGATTGAAAA	12780
ATCATAGACA AAGCGTAGAT AAAGCCGATA ATCAGTGCAA TAACCAAACC ACCAAGTCCA	12840
GATCTTATAA AGAGATAACC AACCGCATAA CCAACAAGAG CTAAGAGTAG GAAAAATACC	12900
AGCAACAAAA TCCAGGTTTT TCGTTTATTG CTTGCAATTG GATCAAACAA CATCTTAGTC	12960
ACCTAAACCG CTAAAATCAA CTTTAGGAAC CGACTTTCC TCTTCAGGTG TTTGAAGGAA	13020
ATCTGCCGCT TTAAATCCAA ACATTCCAGC GATAATATTG CTCGGGAAAG TTTCTAATTG	13080
TACATTGTAG TTGCTGACAA CACTGTTATA GAGTTGACGA GAGTAAGAAA TTTTATTTC	13140
TGTGTTTGTC AACTCCTCTT GCAATTAAAC AAAGTTAGCA CTAGCTTCA AATCTGGATA	13200
GCTTTCTGCA ACTGCAAAAA TACCTGAAAC CTGACGAGTG AGGGCATCAC TGGCTTCAT	13260
AGCTTCTGCT GGTGAAGTCG CTGCCGCCAC TTGGTTACGT AGTTCTGCCA CCTTTCAAG	13320
GGTAGAACCT TCATATTG CATAACCTTT TACAGTCTCA ATCAAGTTTG GCAAGAGGTC	13380
ATTGCGACGT TTCAACTGAA CATCAATCTG ACTCCAAGCC TCCTTGGTT GCATACGATT	13440
TTTAACCAAA CCGTTATAGC TAACAATCAC AAAAATAACA ATAAGAGCGA TAACTCCAAG	13500
AATAATCCAA GTCATAATAT AAGTCCTTTC TGCTTTAGA TTAGTACCAAG TATATCAAAT	13560
TTTCTATGAT TGTGGAAAAA TAAGATGATA CTAAAGAAGG AAATAACTAT GAAACCAAAA	13620
ACATTTTACA ACTTGCTTGC CGAGCAGAAT CTTCCACTTT CGGACCAGCA AAAAGAACAA	13680
TTTGACGTT ATTTTGAGCT CTTGGTCGAG TGGAAATGAGA AGATTAATTG GACGGCGATT	13740
ACGGACAAAGG AAGAAGTTA TCTAAACAT TTTTACGATT CGATTGCACC CATTCTCAA	13800
GGTTTGATTC CCAATGAAAC TATCAAACCTT CTTGATATCG GGGCTGGGC AGGATTTCT	13860
AGTCTACCAA TGAAAATTCT CTATCCGGAG TTAGATGTGA CCATTATTGA TTCACTCAAT	13920
AAGCGCATCA ACTTCTTACA ACTCTGGCT CAAGAACTGG ATTGAAACGG AGTCATTTC	13980
TACCACGGAC GTGCCGAAGA TTTGCCAA GACAAGAACT TCCGTGCTCA ATATGATT	14040
GTAACAGCTC GTGCGGTTGC CCGTATGCAG GTCCTATCTG AATTGACTAT TCCCTACCTT	14100
AAGGTTGGTG GCAAACATTAG AGCACTCAAG GCTAGCAATG CGCCTGAGGA ATTATTAGAA	14160
GCTAAGAATG CCCTCAATCT CCTTTTTAGT AAGGTCGAAG ACAATCTCAG TACGCCCTAC	14220
CGAATAGAGA TCCGCGCTAT ATCACAGTGG TAGAAAAAGAA AAAAGAAACAA CCAAATAAAT	14280
ATCCACGTAA GGCTGGTATG CCAAATAAAC GCCCACTTTA AATTAGTAG TAAACAAATG	14340
TTTACAAAAT CAGCCTCGCT CTTTATTTC TAGGCTCGGG AAAAATGAT TTACAAAATC	14400
AGCCTCGCTC TTTTATTTC AGGCTCGGAA AAAAATGATT TACAAAATCA TTTTTCTG	14460

CTATACTATC CTAAGCAAAG GTTTTTAATG TCATCCCGTG AGGTGACGAA GACGCAGAAA TATTAAAAC TCTTTAAAAT CTAAATTTA AAGAAGTCTT ACTCTGAGGG CCTATTGCTG TAAAATAATG GGCTCTTTT TGATGCCAA AAGTGAGGTT TATATGAAAC AAGAACAC TGTTGATTG TTAC	790 14520 14580 14640 14654
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(2) INFORMATION FOR SEQ ID NO: 107:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6405 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 107:

AGAAAAATCT GCTTACAGA AAATAAAAAT AATAGGAGAA AATCTATGTC AGATTGAAA AAATACGAAG GTGTCATTCC AGCCTCTAC GCATGTTATG ATGATCAAGG AGAAGTAAGC CCAGAACGTA CGCGTGCCTT GGTTCAATAC TTCATTGATA AAGGTGTTCA AGGTCTTTAT GTCAATGGTT CTTCTGGTGA ATGTATCTAC CAAAGCGTTG AAGATCGAA CTTGATTTG GAAGAAGTCA TGGCGGTAGC AAAGGTAAT TGACCATTAT TGCCCATGTT GCTTGCAATA ATACTAAAGA TAGTATGGAA CTTGCTCGCC ATGCTGAAAG CTTGGGAGTA GATGCTATTG CAACGATTCC ACCAATTAT TTCCGCTTGC CAGAATACTC AGTTGCCAAA TACTGGAACG ATATCAGTTC TGCAAGCTCCA AACACAGACT ACGTGATTCA CAACATTCCCT CAATTGGCAG GGGTTGCTTT GACTCCAAGC CTTTACACAG AAATGTTGAA AAATCCTCGT GTTATCGGTG TGAAGAACTC TTCTATGCCA GTTCAAGATA TCCAAACCTT TGTCAGCCTT GGTGGAGAAG ACCATATCGT CTTAATGGT CCTGATGAGC AGTTCCCTAGG AGGACGCCCTC ATGGGGGCTA GGGCTGGTAT CGGTGGTACT TATGGTGCTA TGCCAGAACT CTTCTTGAAA CTCATCAGT TGATTGCGGA TAAGGACCTA GAAACAGCGC GTGAATTGCA GTATGCTATC AACGCAATCA TTGGTAAACT CACTTCTGCT CATGGAAATA TGTACGGTGT CATCAAAGAA GTCTTGAAAA TCAATGAAGG CTTGAATATT GGATCTGTTG GTTCACCATT GACACCAGTG ACTGAAGAAG ATCGTCCAGT TGTAGAAGCG GCTGCTGCCT TGATTCGTGA AACCAAGGAG CGCTTCCTCT AATCTAAAAG GAGGTATTTA TGACATATTA CGTTGCAATT GATATCGGTG GAACCAACAT CAAGTATGGT TTGGTTGATC AAGAGGGCA ACTTCTTGAA TCGCATGAAA TGCCAACTGA GGCGCATAAG GGTGGACCTC ATATCTTACA AAAGACCAAA GATATCGTAG CTAGTTATTT AGAAAAAGGC CCAGTAGCAG GTGTTGCCAT ATCTTCTGCT GGGATGGTGG ATCCGGATAA	60 120 180 240 300 360 420 480 540 600 660 720 780 840 900 960 1020 1080 1140 1200
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GGGTGAGATT TTCTATGCTG GGCGCAAAT CCCTAACTAC GCAGGCACCC AGTTCAAAA	1260
GGAAATCGAA GAAAGCTTTA CTATTCCTTG TGAGATTGAA AATGATGTCA ACTGTGCAGG	1320
TCTTGCTGAG GCAGTATCTG GTTCAGGCAA GGGAGCAAGT GTGACACTTT GCTTGACCAT	1380
TGGAACCGGT ATCGGTGGTT GCTTGATTAT GGATAGGAAA GTCTTCCATG GTTTTAGCAA	1440
TTCAGCCTGT GAAGTCGGGT ATATGCATAT GCAGGATGGA GCTTTCAAG ACTTGGCTTC	1500
TACAACAGCT TTAGTGAAT ATGTAGCTGA AGCCCATGGA GAAGATGTTG ATCAGTGGAA	1560
TGGCCGTAGA ATTTCAAAG AAGCCACTGA AGGAAACAAA ATCTGCATGG AAGGTATTGA	1620
CCGTATGGTT GACTATCTAG GAAAAGGTCT GGCAAATATT TGCTACGTTG CCAATCCAGA	1680
AGTGGTTATT CTTGGTGGTG GTATCATGGG GCAAGAGGCT ATCCTCAAAC CTAAGATCCG	1740
TACAGCCTTG AAAGAGGCTT TGGTACCAAG TTTAGCAGAA AAAACACGAT TAGAATTGCG	1800
CCATCACCAA AATACAGCAG GGATGTTGGG TGCAATTATT CATTAAAGA CAAACAAATC	1860
CTAGTTTGGC TCAGCCAAAC TAGGATTTTC TTACACGTTT TTGTCTACGA TAGCCGTTGA	1920
GTTTTTATT TTCCCAGTAG CTATTAAAGA TTTTTCCCTT GCTTCGCGA TTGATTCCA	1980
AAAAGTAGGC ATAATCAAA TCGATAAAGA AGAGCATAGG AAGTTGAGCG GATATTGTT	2040
GGATATAGGA GGGTTGGCTG TGGGTTGGCTA CAAGAACAGT CTCTGTATAG GTCTGGCTAT	2100
CTTTATTGGG AACACTTGTA AAGAGTACAG TCTTGTCCCC CATCTCCTTA GCATCTAATA	2160
GACTATCTAA AATAGAAGGA GTTGAGCCTG AAAGTGAGAA GCCCAGTAAGT AGACAATTTC	2220
CATCCATGAT GCTGGTTGTC CAGGCAAAGC CGTCTGGTC TGTCAGACT TCGCAGACCA	2280
CACCTAGTCG CATAAAACGT AATTTCATTT CACGGGCGAC GAGGCCAGAA CTCCCTGTT	2340
CAAAGAAGTA GATACGCTCA GCATCTCGA TTAGCTGGC AATTGTTCT AGTTGGATT	2400
CGTCAATCAA GTCTTGTT TGTTCCCTCA TATTGCTATA ACTTCTGAGG ACTCGTTGG	2460
TCAGTGGACT GTGCTTGGAG ACTTGGTTGG CTTGATTTTC TGCTGATGT TGGTATTGGA	2520
AAATAAATTC TCGGTAGCCA GTAAAGCCAC ACTTTTAGC AAAGCGGGTC AAAGCAGCTT	2580
GAGAAATATG TAATTTTGG GTGACTTGTT GAGAAGATAA ATCATCTGTA ATCGTTTCAG	2640
CTTGCAAAAA ATAGCGAGCG ATTTCTTGTT CTAGGTCTGT CATTCTCTCA AAATGTGAAT	2700
CAATGATAGT TGCGATATCT GGTTTGTCCA TAGGGAAAGC TCCTTACAT GAGTCATACT	2760
GGAAGACTAG ATCAGAGAAT AGTCACACTT CATTATAACA CATAATATAA GGATAGATAA	2820
ATAAAAACGC ATCTCTGTT TAAAAACGAA AAAATCGAAA AAGCTTCTCT CTTTCCATA	2880
ATTTTCTACT CAAATTGTGG TACAATTAAG AGTAAGATT TAAGTTAGAA ATGAGACTGA	2940

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TTTGTATGAG AAAATTTAAC AGCCATTGCA TTCCGATTG GCTTAATTAA TTGTTCGAA	3000
TCGTCATTG ACTCTTTATG ACCATTATTG GTCGTTGTT GTATATGCAG GTTTGAACA	3060
AGGATTTTA CGAAAAAAAG CTAGCTTCAG CTAGTCAGAC CAAGATTACA AGCAGTTCA	3120
CCCGTGGGA AATTATGAT GCTAGTGGAA AACCTTGTT AGAAAATACG TTAAAGCAGG	3180
TTGTTCCCTT TACCGTAGC AATAAAATGA CGGCTACAGA CTTAAAAGAA ACAGCTAAA	3240
AGTTACTGAC TTATGTGAGC ATCAGTTCTC CAAATTGAC AGAACGCCAG CTGGCGGATT	3300
ACTATTTGGC TGATCCTGAA ATCTATAAAA AAATAGTGGA AGCTCTCCCA AGTGAGAAC	3360
GCTTGGATTC AGATGCAAT CGTCTATCCG AATCAGAACT GTATAACAAT GCGGTCGATA	3420
GTGTACAAC GAGTCAACTA AACTATACAG AGGATGAAA GAAAGAAATC TATCTTTTA	3480
GTCAGTTAAA TGCTGTTGGA AACCTTGCGA CAGGAACCAT TGCGACAGAT CCTCTAAATG	3540
ATTCTCAGGT GGCTGTTATT GCCTCTATTT CAAAGGAGAT GCCTGGCATT AGTATTCTA	3600
CTTCTTGGGA TAGAAAGGTT TTGGAAACTT CCCTTCTTC TATAGTTGG AGTGTATCCA	3660
GTAAAAAGC TGGTCTCCCA GCGGAAGAAG CAGAAGCCTA TCTTAAAAAA GGCTATTCTC	3720
TAAATGACCG TGTAGGAACC TCCTATTTGG AAAAGCAATA TGAAGAGACC TTACAAGGA	3780
AACGCTCGGT AAAAGAAATC CATCTGGATA AATATGGCAA TATGGAAAGC GTGGATACAA	3840
TTGAGGAAGG TAGTAAGGGA AACAAATATCA AACTGACCCT TGATTTGGCT TTCCAAGATA	3900
GCGTGGATGC TTTACTGAA AGTTATTTCA ATTCTGAGCT AGAAAATGGT GGAGCCAAGT	3960
ATTCTGAAGG TGTCTATGCA GTCGCCCTTA ACCCAAAAC AGGTGCGGTT TTGTCTATGT	4020
CAGGGATTAA ACATGACTTG AAAACGGGAG AGTTGACGCC TGATTCCTTG GGAACGGTAA	4080
CCAATGTCCTT TGTTCCAGGT TCGGTTGTCA AGGCGGCGAC CATCAGCTCA GGTTGGAAA	4140
ATGGAGTCCTT GTCAGGAAAC CAGACCTTGA CAGACCAGTC CATTGTCCTTC CAAGGTTCA	4200
CTCCCACCAA TTCTGGTAT ACTCAGGCTT ACGGTTCAATT CCCTATCACA GCGGTCCTAAG	4260
CTCTGGAGTA TTCATCAAAT ACCTATATGG TCCAAACAGC CTTAGGTCTT ATGGGGCAA	4320
CCTATCAACC CAATATGTTT GTCGGCACCA GCAATCTAGA GTCTGCTATG GAGAAACTGC	4380
GTTCAACCTT TGGCGAATAT GGCTTGGGTA CTGCGACAGG AATTGACCTA CCAGATGAAT	4440
CTACTGGATT TGTTCCAAA GAGTATAGCT TTGCTAATTA CATTACTAAT GCCTTGGGC	4500
AGTTTGATAA CTATACGCCG ATGCAGTTGG CTCAGTATGT AGCAACTATT GCAAATAATG	4560
GTGTTCGTGT GGCTCCTCGT ATTGTTGAAG GCATTTATGG TAATAATGAT AAGGGAGGAC	4620
TGGGTGACTT GATTCAAGCAA CTGCAACCGA CAGAGATGAA TAAGGTCAAT ATATCCGACT	4680
CCGATATGAG CATCTTGCAC CAAGGTTTTT ATCAGGTTGC CCATGGTACT AGTGGATTGA	4740

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CAACTGGACG	TGCCTTTCA	AATGGTGCCT	TGGTATCCAT	TAGCGAAAAA	ACAGGTACAG	4800
CCGAAAGCTA	TGTGGCAGAT	GGTCAGCAAG	CAACCAATAC	CAATGCGGTG	GCCTATGCC	4860
CATCTGATAA	TCCCCAAATC	GCTGTCGAG	TGGTCTTCC	TCATAATACC	AATCTAACAA	4920
ATGGTGTAGG	ACCTTCCATT	GCGCGTGACA	TTATCAATCT	GTATCAAAAAA	TACCATCCAA	4980
TGAATTAGAA	AGGAATTAT	GCTTTATCCA	ACACCTATTG	CCAAGTTGAT	TGACAGTTAT	5040
TCTAAGTTAC	CAGGTATCGG	GATTAAGACG	CCTACGCGTC	TGGCCTTTA	TACGATTGGG	5100
ATGTCTGCTG	ATGATGCAA	TGAATTGCA	AAAATCTCC	TTTCTGCTAA	GAGAGAATTG	5160
ACATATTGTT	CTATTTGAG	ACGTTTGACA	GACGACGATC	CTTGTCTAT	CTGTACTGAT	5220
CCGACTCGTG	ACCAGACAAC	AATTTAGTT	CTTGAGGATA	GTAGAGATGT	GGCAGCCATG	5280
GAAAATATCC	AAGAATACCA	TGGACTCTAT	CATGTCCTTC	ATGGCCTCAT	TTCTCCTATG	5340
AATGGTATCA	GTCCGGACGA	TATCAATCTC	AAGAGCCTTA	TGACTCGTCT	TATGGATAGT	5400
GAGGTTTCAG	AAGTGATTGT	GGCGACTAAT	GCTACAGCGG	ATGGTGAAGC	GACTTCCATG	5460
TATCTTCAC	GTGGCTCAA	GCCGGCTGGT	ATCAAGGTTA	CGCGTCTAGC	ACGAGGTCTC	5520
GCTGTGGGAG	CGGACATTGA	GTATGCGGAC	GAAGTGACAC	TCTTACGAGC	CATTGAAAAT	5580
CGGACAGAGT	TGTAAGTGT	GGCAAATTAA	CGAACTCCAT	TCATTTATAA	AAAATCAAAG	5640
AGGCTGAAA	TCGTTCTAT	CGGCCTCTTT	TTGTATAGTG	TGATGAGTAG	GCTCAGGTTC	5700
AAGTTTAAA	AAACCAAGCA	AATATGATAT	ACTAAAGAGC	GAGTATTCTA	GTAGAATTAG	5760
GACAAATAAT	ATGAAACAAA	CGATTATTCT	TTTATATGGT	GGACGGAGTG	CGGAACGCGA	5820
AGTCTCTGTC	CTTTCAGCTG	AGAGTGTAT	CGGTGCGGTG	GATTACGACC	GTTCACAGT	5880
CAAGACTTTC	TTTATCAGTC	AGTCAGGTGA	CTTTATCAA	ACACAGGAAT	TTAGTCATGC	5940
TCCGGGGCAA	GAAGACCGTC	TCATGACCAA	TGAAACCATT	GATTGGGATA	AGAAAGTTGC	6000
ACCAAGTGCT	ATCTACGAAG	AAGGTGCAGT	GGTCTTCCA	GTCCTTCACG	GGCCAATGGG	6060
AGAAGATGGC	TCTGTTCAAG	GATTCTTGG	AGTTTGAAA	ATGCCTTACG	TTGGTTGCAA	6120
CATTTGTCA	TCAAGTCTT	CCATGGATAA	AATCACGACT	AAGCGTGTTC	TGGAATCTGC	6180
TGGTATTGCC	CAAGTCCCTT	ATGTGGCTAT	CGTTGAAGGC	GATGATGTGA	CTGCTAAAAT	6240
CGCTGAAGTG	GAAGAAAAAT	TGGCTTATCC	AGTCTTCACT	AAGCCGTCAA	ACATGGGGTC	6300
TAGTGTGGT	ATTTCTAAGT	CTGAAAACCA	AGAAGAACTC	CGTCAAGCCT	TAAAACATTGC	6360
CTTCCGATAT	GACAGCCGTG	TCTTGGTTGA	GCAAGGAGTG	AATGC		6405

(2) INFORMATION FOR SEQ ID NO: 108:

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(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 11309 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 108:

CGAGCTCGGG TACCGGGATT TTAAGGAGTT TGATATGTAT AACCTATTAT TAACCATT	60
ATTAGTATTA TCTGTTGTGA TTGTGATTGC AATTTCATG CAACCAACCA AAAACCAATC	120
CAGCAATGTA TTTGATGCCA GTTCAGGTGA TTTGTTGAA CGCAGTAAAG CTCGCGTTT	180
TGAAGCTGTA ATGCACCGTT TGACAGGGAT TTTAGTCTTT TTCTGGCTAG CCATTGCCTT	240
ACCATTGACG GTATTATCAA GTAGATAAGA AAATAATGGG CAGGACTAGG TCTTTGCCTC	300
TTTTTATTGT TAAAGGATGT TTGAGAAGGT TTTACAGTAA AAGAAAATTA AAAAATCTAG	360
AAAGAAAATA TGAAAGATAG AATAAAAGAA TATTTACAAG ACAAGGGAAA GGTGACTGTT	420
AATGATTGGG CTCAGGCTTT GGGAAAAGAC AGTTCCAAGG ATTTTCGTGA GTTGATTAAA	480
ACCTTGTCCCT TAATGGAAAG AAAGCACCAA ATTCTGTTTG AAGAAGATGG TAGTCTGACA	540
TTAGAAAATTA AGAAAAAACAA TGAGATTACC CTCAAGGGGA TTTTTCATGC CCATAAAAAT	600
GGCTTTGGCT TTGTTAGTCT GGAAGGCGAG GAGGACGACC TTTTTGTAGG GAAAATGAT	660
GTCAACTATG CTATTGATGG TGATACCGTC GAGGTAGTGA TTAAGAAAGT CGCTGACCGC	720
AATAAGGGAA CAGCAGCAGA AGCCAAAATT ATTGATATCC TAGAACACAG TTTGACAACA	780
GTTGTCGGGC AAATCGTTCT GGATCAGGAA AAACCTAAGT ATGCTGGCTA TATTGTTCA	840
AAAATCAGA AAATCAGTCA ACCGATTAT GTTAAGAAC CAGCCCTAAA ATTAGAAGGA	900
ACAGAAGTTC TCAAAGTCTT TATCGATAAA TACCCAAGCA AGAAACATGA TTTCTTGTC	960
CGCAGTGTTC TCGATGTAGT GGGACACTCA ACGGATGTGCG GAATTGATGT TCTTGAGGTC	1020
TTGGAATCAA TGGACATTGT ATCCGAGTTT CCAGAAGCTG TTGTTAAGGA AGCAGAAAGT	1080
GTCGCTGATG CTCCGCTCTCA AAAGGATATG GAAGGTCGTC TGGATCTAAG AGATGAAATT	1140
ACCTTTACCA TTGACGGTGC GGATGCCAAG GACTTGGACCG ATGCAGTGCA TATCAAGGCT	1200
CTGAAAATG GCAATCTGGA GTTGGGGTT CACATCGCAG ATGTTCTTA TTATGTGACC	1260
GAGGGGTCTG CCCTGACAA GGAAGCCCTT AACCGTGCAG CTTCTGTTA CGTGACAGAC	1320
CGAGTGGTGC CAATGCTTCC AGAACGACTA TCAAATGGCA TCTGCTCTCT CAATCCCCAA	1380
GTTGACCGCC TGACCCAGTC TGCTATTATG GAGATTGATA AACATGGTCG TGTGGTCAAC	1440
TATACCATTA CACAAACAGT TATCAAGACC AGTTTCGTA TGACCTATAG CGATGTCAAT	1500

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GATATCCTAG	CTGGCGATGA	AGAAAAGAGA	AAAGAATATC	ATAAAATTGT	ATCAAGTATC	1560
GAACTCATGG	CCAAGCTTCA	TGAAACTTTA	GAAAACATGC	GTGTGAAACG	TGGAGCTCTC	1620
AATTTGATA	CCAATGAAGC	GAAGATTTA	GTGGATAAAC	AAGGTAAGCC	TGTTGATATC	1680
GTTCTTCGGC	AGCGTGGTAT	TGCCGAGCGG	ATGATTGAGT	CTTTTATGTT	GATGGCTAAT	1740
GAAACAGTTG	CCGAACATTT	CAGCAAGTTG	GATTTGCCTT	TTATCTATCG	AATTCAACGAG	1800
GAGCCTAAGG	CTGAAAAGGT	TCAGAAGTTT	ATTGATTATG	CTTCGAGTTT	TGGCTTGC	1860
ATTTATGGAA	CTGCCAGTGA	GATTAGTCAG	GAGGCACCTTC	AAGACATCAT	GCGTGCTGTT	1920
GAGGGAGAAC	CTTATGCAGA	TGTATTGTCC	ATGATGCTTC	TCGCTCTAT	GCAGCAGGCT	1980
CGTTATTTCGG	AGCACAAATCA	CGGCCACTAT	GGACTAGCTG	CTGACTATTA	TACTCACTTT	2040
ACCAGTCAA	TTCGTCGTTA	TCCAGACCTT	CTTGTTCAACC	GTATGATTG	GGATTACGGC	2100
CGTTCTAAGG	AAATAGCAGA	GCATTTGAA	CAACTGATTTC	CAGAGATTG	GACCCAGTCT	2160
TCCAACCGTG	AACGTCGTGC	CATAGAAGCT	GAGCGTGAAG	TCGAAGCCAT	GAAAAAGGCT	2220
GAGTATATGG	AAGAATACGT	GGGTGAAGAG	TATGATGCAG	TTGTATCAAG	TATTGTCAAA	2280
TTCCGGTCTCT	TTGTCGAATT	GCCAAACACA	GTTGAAGGCT	TGATTCACAT	CACTAATCTG	2340
CCTGAATTTC	ATCATTCAA	TGAGCGTGAT	TTGACTCTTC	GTGGAGAAAA	ATCAGGTATC	2400
ACTTTCCGAG	TGGGTCAAGCA	GATCCGTATC	CGTGTGAAA	GAGCGGATAA	AATGACTGGA	2460
GAGATTGATT	TTTCATTCTG	ACCTAGTGAG	TTTGATGTGA	TTGAAAAAAGG	CTTGAAACAG	2520
TCTAGTCGTA	GTGGCAGAGG	GCGTGATTCA	AATCGTCGTT	CGGATAAGAA	GGAAGACAAG	2580
AGAAAATCAG	GACGCTAAA	TGATAAGCGT	AAGCATTAC	AAAAAGACAA	GAAGAAAAAA	2640
GGAAAGAAAC	CTTTTTACAA	GGAAAGTAGCT	AAGAAAGGAG	CCAAGCATGG	CAAAGGGCGA	2700
GGGAAAGGTC	GTCGCACAAA	ATAAAAGGC	ACGCCACGAC	TATACAATCG	TAGATACGCT	2760
AGAGGCAGGG	ATGGTCTCTGA	CTGGAACTGA	AATCAAGAGT	GTACGAGCTG	CTCGAATTAA	2820
TCTCAAGGAT	GGCTTGCTC	AAGTAAAAAA	TGGAGAAGTT	TGGCTGAGCA	ATGTTCATAT	2880
CGCGCCTTAC	GAAGAGGGCA	ATATCTGGAA	CCAGGAACCA	GAACGTCGTC	GTAAACTCCT	2940
GCTCCATAAA	AAGCAAATTC	AAAAATTGGA	ACAAGAGATC	AAAGGGACAG	GAATGACCTT	3000
AGTTCCCCCTT	AAGGTCTATA	AAAAAGATGG	CTACGCTAAG	CTTCTTTAG	GACTTGCCAA	3060
AGGGAAGCAT	GACTATGACA	AACGGGAGTC	TATCAAACGT	CGTGAGAAA	ATCGAGATAT	3120
CGCGCGTGTG	ATGAAAGCTG	TTAACAGCG	ATAAAAAGAG	GAATTGAAAA	TGGAAAATT	3180
AGTTGCCTAT	AAACGCATGC	CTTTGTGGAA	TAAACAAACAA	ATGCCTGAAG	CTGTTCAAGCA	3240

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AAAGCACAAT	ACAAAAGTTG	GGACTTGGGG	GAAAATTACT	GTCTTGAAGG	GAGCTCTCAA	3300
GTTTATTGAA	TTGACAGAAG	AAGGGGAAGT	TCTAGCTGAA	CACCTCTTG	AAGCAGGGC	3360
AGACAATCCA	ATGGCCCAAC	CTCAAGCCTG	GCACCGAGTG	GAAGCTGCCA	CAGATGATGT	3420
GGAATGGTAC	TTGGAATTTC	ATTGTAACC	TGAGGATTAT	TTTGCTAAAA	AATACAATAC	3480
CAATCCTGTT	CATTCAGAGG	TCCTAGAGGC	CATGCAGACA	GTGAAACAAG	GGAAAGCTTT	3540
GGATTTGGGT	TGTGGTCAGG	GGCGTAATTC	TCTTTTCTA	GCCCAGCAAG	ATTTTGATGT	3600
GACGGCTGTA	GATCAAAATG	GACTAGCTCT	TGAAATCTTG	CAAAGCATTG	TGGAGCAGGA	3660
AGATTTGGAC	ATGCCTGTTG	GCCTTACGA	TATCAATTCA	GCTAGCATTG	AACAAGAATA	3720
TGATTTTATC	GTTCACACAG	TTGTTCTCAT	GTTCCTACAA	GCGGACCGCA	TTCCAGCTAT	3780
TATTCAAAAT	ATGCAGGAGA	AAACCAAGTGT	TGGTGGTTAC	AACCTTATCG	TTTGTGCCAT	3840
GGACACGGAG	GATTATCCTT	GCTCGGTTAA	CTTCCCATTG	ACCTTTAAAG	AAGGAGAACT	3900
GGCAGACTAT	TACAAGGATT	GGGAATTGGT	TAAGTACAAT	GAAAATCCAG	GCCATTTGCA	3960
CCGTCGCGAT	GAGAATGGCA	ATCGTATTCA	ACTACGCTTT	GCGACCTTAC	TAGCTAAGAA	4020
AATCAAGTAA	ACACACATGA	AGATTAGGAA	TTTTCTGAT	CTTTTTCTT	TTTTACGAAT	4080
GATATAGAAA	AGGAGGGAAT	TCATGTTGT	TGCGAGAGAT	GCTAGGGAG	AATTGGTAAA	4140
TGTGTTAGAG	GATAAACTTG	AGAACGAAAGC	ATACACCTGC	CCAGCTGTG	GAGGCCAGCT	4200
CCATTTGCGT	CAAGGACCAA	GTGTACGGAC	GCATTTGCC	CATAAATCCT	AAAAAGACTG	4260
TGATTTTTTC	TTTGAAAATG	AAAGTCCAGA	ACACCTGGCC	AATAAGGAAT	CCCTCTATCA	4320
CTGGTTGAAA	AAAGAGACAA	AGGTTCAATT	AGAGTACCCG	CTTTCAGAAC	TTAAACAGAT	4380
TGCGGATGTA	TTTGAAATG	GCAATCTAGC	TCTAGAAGTT	CAGTGTAGTC	CCTTGCCTCA	4440
GAAAGTCCTT	AAAGAGCGAA	GTGAGGGCTA	TCGTAGTCAG	GGTTACCAAG	TACTGTGGTT	4500
GCTGGGTCAA	AAACTGTGGC	TCAAGGAGCG	TTTGACTCGT	CTACAGCAAG	GTTTCTTTA	4560
TTTCAGTCAA	AACATGGGCT	TTTATGTTG	GGAATTAGAC	AAGGAAAAAC	AAGTTTTAAG	4620
ACTCAAATAC	CTGATTTACC	AGGATCTCCG	CGGTAAACTC	CATTATCAA	TCAAGGAATT	4680
TTCCTATGGT	CAAGGTAGTT	TATTGGAAAT	ATTGCGTCTT	CCCTATAAGA	GACAAAAAAT	4740
ATCTCATTTC	ACAGTTCTG	AGGACAAGGA	CATCTGTCGC	TATATCCGGC	AACAACCTTA	4800
TTATCAAAAT	CTCTTTGGA	TGAAAGAAC	AGCAGAAGCC	TATCAAAGG	GAGAAAATAT	4860
CCTGACTTAT	GGACTGAAAG	AATGGTATCC	ACAAATTGCA	CCAATAGTGG	GCAAATTTT	4920
CCAGATTGAA	CAAGACTTGA	CTAGCTATTA	TCAGCACTTT	TATACCTATT	ACCAAAAAAA	4980
TCCTCAAAAT	GATTGGAAA	AGCTTTATCC	ACCAGCCTTT	TATCAGCAAT	ATTTCTTGAA	5040

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AAATATGGTA	GAATAGAAAAG	GATGGAGGAA	TCTAATGGTA	TTACAAAGAA	ATGAAATAAA	5100
TGAAAAAGAT	ACATGGGATC	TATCAACGAT	CTACCCAACT	GACCAGGCTT	GGGAAGAAC	5160
CTTAAAAGAT	TTAACAGAAC	AATTGGAGAC	AGTAGCCAG	TATGAAGGCC	ATCTCTTGGA	5220
TAGTGCGGAT	AACCTACTAG	AAATCACTGA	ATTTTCTCTT	GAAATGGAAC	GCCAGATAGA	5280
GAAGCTTAC	GCTTATGCTC	ATATGAAGAA	TGACCAGGAT	ACACGTGAAG	CTAAGTATCA	5340
AGAGTACTAT	GCCAAGGCCA	TGACACTCTA	CAGCCAGTTA	GACCAAGCCT	TTTCATTCTA	5400
TGAGCCTGAA	TTTATGGAGA	TTAGCGAAAA	GCAGTATGCT	GACTTTTAG	AAGCTCAACC	5460
AAAGCTGCAG	GTTTATCAAC	ACTATTTGA	CAAGCTTTG	CAAGGCAAGG	ATCACGTTCT	5520
TTCACAACGT	GAAGAAGAAT	TATTGGCTGG	AGCTGGAGAA	ATCTTGGTT	CAGCAAGTGA	5580
AACCTTCGCT	ATCTTGGACA	ATGCGGATAT	TGTGTTCCCT	TATGTCTAG	ACGATGATGG	5640
TAAAGAACGT	CAGCTATCTC	ATGGGACTTA	CACACGTTG	ATGGAGTCTA	AAAACGTGA	5700
GGTTCGCCGT	GGTGCCCTATC	AAGCTTTA	TGCGACTTAC	GAACAATTCC	AACACACCTA	5760
TGCCAAAACC	TTGCAAACCA	ATGTTAAGGT	GCAAAATTAC	CGTGCTAAAG	TTCGTAACTA	5820
CAAGAGTGCT	CGTCATGCAG	CCCTCGCAGC	GAATTTGTT	CCAGAAAGTG	TTTATGACAA	5880
TTTGGTAGCA	GCAGTTCGCA	AGCATTGCCC	ACTCTTACAT	CGCTATCTG	AGCTTCGTT	5940
AAAAATCTTG	GGGATTCAG	ATCTCAAGAT	GTACGATGTC	TACACACCGC	TTTCATCTGT	6000
TGAATACAGT	TTTACCTACC	AAGAAGCCTT	GAAAAAAAGCA	GAAGATGCTT	TGGCAGTCTT	6060
GGGTGAGGAT	TACTTGAGCC	GTGTTAACG	TGCCTTCAGC	GAGCGTTGGA	TTGATGTTA	6120
CGAAAATCAA	GGCAAGCGTT	CAGGTGCCTA	CTCTGGTGGT	TCTTATGATA	CCAATGCCTT	6180
TATGCTTCTC	AACTGGCAAG	ACAATCTGGA	CAATCTCTTT	ACTCTGTTC	ATGAAACAGG	6240
TCACAGTATG	CATTCAAGCT	ATACTCGTGA	AACTCAGCCT	TATGTTACG	GGGATTACTC	6300
TATCTTTTG	GCTGAGATTG	CCTCAACTAC	CAATGAAAAT	ATCTTGACGG	AGAAATTATT	6360
GGAAGAAGTG	GAAGACGACG	CAACACGCTT	TGCTATTCTC	AATAACTTCC	TAGATGGTT	6420
CCGTGGAACA	GTTCGGGCC	AAACTCAATT	TGCTGAGTT	GAACACGCCA	TTCACCAAGC	6480
AGATCAAAAT	GGGGAGGTCT	TGACAAGCGA	TTTCCTAAAT	AAACTCTACG	CAGACTTGAA	6540
CCAAGAGTAT	TATGGTTTGA	GTAAGGAAGA	CAATCCTGAA	ATCCAATACG	AGTGGGCTCG	6600
CATTCCACAC	TTCTACTATA	ACTACTATGT	ATATCAATAT	TCAACTGGCT	TTGGGGCCGC	6660
CTCAGCCTG	GCTGAAAAAA	TTGTCCATGG	TAGTCAAGAA	GACCGTGACC	GCTATATCGA	6720
CTACCTCAAG	GCAGGTAAGT	CGGACTATCC	ACTTAATGTC	ATGAGAAAAG	CTGGTGTTGA	6780

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TATGGAGAAG	GAAGACTACC	TCAACGATGC	CTTTGCAGTC	TTTGAACGCC	GTTTAAATGA	6840
GTTTGAAGCC	CTTGTGAAA	AATTAGGATT	GGCATAAAAT	GGTTGAATCG	TATAGTAAGA	6900
ATGCTAACCA	TAACATGCGT	CGTCCTGTCG	TCAAAGAAGA	AATTGTAGAC	TTGATGCGTC	6960
AGCGTCAAAA	GCAGGTCACA	GGTTCTTGA	AAGAATTGGA	AGACTTGC	CGCAAGGAAA	7020
ATATTCCAT	TATTCCCCAT	GAAACGGTTG	CTTATTTCCG	TTTCTTATG	GAAACCATGC	7080
AGCCTAAAAA	TATTCTGGAA	ATTGGGACGG	CTATCGTTT	TCAGCTCTC	TTGATGGCTG	7140
AACATGCGCC	AAATGCTAAG	ATTACAACTA	TTGATCGTAA	TCCAGAAATG	ATTGGTTTG	7200
CCAAGGAAAA	TTTTGCCAG	TTTGACAGTC	GCAAGCAAAT	CACTCCCTA	GAGGGAGATG	7260
CGGTGGATGT	CTTATCTACA	CTGACAGAGT	CTTATGATTT	CGTCTTATG	GATTCTGCCA	7320
AGTCTAAATA	CATCGTCTT	CTGCCAGAAA	TCCTCAAACA	TTTGGAAAGTT	GGTGGTGTGG	7380
TTGTCTTGG	TGATATTTT	CAAGGTGGT	ATGTTGCCAA	GGATATTATG	GAAGTCCGTC	7440
GTGGTCAGCG	AACCATTAT	CGAGGCCTTC	AAAAATTATT	TGATGCAACC	TTAGACAATC	7500
CAGAACTCAC	CGCAACATTA	GTGCCCTTAG	GAGATGGTAT	TCTCATGCTT	CGTAAAAATG	7560
TAGCAGATGT	TCAACTGTCT	GAAAGCGAAT	GATTTTCAGA	AAAATTAAAG	AAAAAATAGT	7620
AAAATAGATA	GAGTAACACT	TATCTCAAAG	GAGTAGACAT	GAAGAAAAAA	TTATTGGCAG	7680
GTGCCATCAC	ACTATTATCA	GTAGCAACTT	TAGCAGCTTG	TTCGAAAGGG	TCAGAAGGTG	7740
CAGACCTTAT	CAGCATGAAA	GGGGATGTCA	TTACAGAACAA	TCAATTAT	GAGCAAGTGA	7800
AAAGCAACCC	TTCAGCCAA	CAAGTCTTGT	TAAATATGAC	CATCCAAAAA	GTTTTGAAA	7860
AACAATATGG	CTCAGAGCTT	GATGATAAAG	AGGTTGATGA	TACTATTGCC	GAAGAAAAAA	7920
AACAATATGG	CGAAAACACTAC	CAACGTGTCT	TGTCACAAGC	AGGTATGACT	CTTGAAACAC	7980
GTAAAGCTCA	AATTCTACA	AGTAAATTAG	TTGAGTTGGC	AGTTAAGAAG	GTAGCAGAAG	8040
CTGAATTGAC	AGATGAAGCC	TATAAGAACAG	CCTTGATGA	GTACACTCCA	GATGTAACGG	8100
CTCAAATCAT	CCGTCTTAAT	AATGAAGATA	AGGCCAAAGA	AGTTCTCGAA	AAAGCCAAGG	8160
CAGAAGGTGC	TGATTTGCT	CAATTAGCCA	AAGATAATTC	AACTGATGAA	AAAACAAAAG	8220
AAAATGGTGG	AGAAATTACC	TTTGATTCTG	CTTCAACAGA	AGTACCTGAG	CAAGTCAAAA	8280
AAGCCGCTT	CGCTTAGAT	GTGGATGGTG	TTTCTGATGT	GATTACAGCA	ACTGGCACAC	8340
AAGCCTACAG	TAGCCAATAT	TACATTGTA	AACTCACTAA	GAAAACAGAA	AAATCATCTA	8400
ATATTGATGA	CTACAAAGAA	AAATTAAAAA	CTGTTATCTT	GAETCAAAAA	CAAAATGATT	8460
CAACATTGT	TCAAAGCATT	ATCGGAAAAG	AATTGCAAGC	AGCCAATATC	AAGGTTAAGG	8520
ACCAAGCCTT	CCAAAATATC	TTTACCCAAT	ATATCGGTGG	TGGAGATTCA	AGCTCAAGCA	8580

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GTAGTACATC AAACGAATAG TCCAAATCAA TGAGTCAGGG AAAAAACTCG ACTTCAGGAA	8640
AAAATGAAGC AAACATTCCC ACAATAAAC GCATAGTACA AGGTTGTAC TGCCCCCAA	8700
AAAGTTAGAC ATTAATTAA TCCGAAGGAT TTAGTTCTGT ATTGCACAGA GCTAAGTCCT	8760
TTTAGTTTA TCTTAATTCT CTTATTGTTG TAATAATCAA TATAGTCTAT AATGGCTCGT	8820
TCCAATTGAT TAAGTGTATT AAAATGTTTC TCATAGCCAT AAAACATTTC GGATTTAAA	8880
ATGCCAAAGA AAGATTCAT CCTACCGTTG TCTTGGCTGT TGCCCTTACG TGACATGGAT	8940
GCTTGAATTC CCTTACTCTC TAGGAAGCGA TGATAAGAAT CGTGTGATA TTGCCAGCCT	9000
TGGTCACTAT GGAGAATCGT ATTCTCGTAG TGCTTCTCTT TGAATGCCGT TTCCAACATT	9060
AACGATCAAT CAATTTAACAT ATGTACCTAA GATTAGAATT GTTTATCCA AATTATTTG	9120
AAAGCTTCTC TAAGCTATAT CCTTGTTTC TAAGTTCATA GATCTGAAC TTATCATCAT	9180
AAGTTAATTT CATAATAAAA ACACCCAAA AGTTAGATT TTTCTGTCTA ACTTTGGGG	9240
TGTAGTTCAT GTACACCTGA TATGATGCGT TTTATAATTT TAAAGACTTT TTGACCAGCC	9300
TCATTTTTT AACTTGATAC TCAGTGAAAA GCAAAGATTA AACTAGGAAG CTAGCTGTAG	9360
GCTGCTCAAA GAACAGCTTT GAGGTTGTAG ATAAAACCTTG TGAGGTCAAC AACATATATA	9420
ATGTGAAGCT GACGTGGTTT GAATAGATT TAGAAGAGTA TGAGTCTGGA AGTTTAATG	9480
GATAATGCAA GATTCATAG AATGGTAAG CTAGAGTTCT TATGTGAAGA GTTGGGCAT	9540
AAACTTTAC CTTTCCCTCC CTACTCATCT TAGTATAGAA AAGTGAATCT GAAATAGTAC	9600
ATAACTGCTT CTAAAACATT CTTATAAATT GATTTAAATT CTCAAATCAT ATTATTCAGT	9660
TCTTATTTC TTTGTTCTA CAATCCTGTT GAGAAGACAC GTGTTCATAT CAAAAAGGTA	9720
TTGGCAAGTT GCAATACCTT TTTACGAGGC TCTGTTGTCT TATTTTGTT TCAACTGACT	9780
ATATCTCCTA TGGTTCTAGT TCAGAAGGCT AGGCTATAAT TATGATTGAT AAGAAGTATC	9840
ATTCCAAGTA TTGGGAGTGA ATGTTTCAAA ATCATGGGTT TCTATAATGG TCAGGCTGGC	9900
ATTTGCTAGA CCGCCATCTT TACGAAGAAG TGGTTCTTTA TAGCCTAGGA GAGTACGAAG	9960
ACTGGCAGTA AGATTGGCGC CGTGTCCGAC AATTAGAATA CGTTCAGCTG GACTATCTTT	10020
TAATGATTG ATAAATTGGA TGGTCCGTTG AGTTGTACTA TAGAGGGATT CGGCTCCGAA	10080
CATTGAGTG TCAAATTGAG CAAGATTGGA ACGAAAAGCC TGGATTGTT GCGGGTAAAT	10140
AGCTTCCAAG GTTGAATTT TCAAACCTTC TAACTTCCCA AGTTGCCATT CACGGAGATT	10200
AGGAACGATT TCTAAAGAAC AGGGGGTATA GAGTTGACTT TGGATAATCT CAGCAGATT	10260
GACCGCTCGA GGTAAATCAC TTGAATAAT CTGATCAAA GGAATTTCCT TGAGATACTG	10320

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ACCAAGTCGT	TTT	AGGGTTT	CAATGGATT	CAGGAAGAAGA	GGAGAAC	TAC	CACTAGCACC	10380
TTGAAAACGA	CCT	TCTGGT	TCCAGAGGGT	ACGACCGTGG	CGGACAAAGT	AGAGTT	CAT	10440
TACTTGATGT	CCT	CCAAAAT	ATCTACAAAG	TCTGCCTTA	CAAAGCTAGC	CAAGT	CTTGT	10500
GGCGCGACGA	TAAT	GCTGTG	TCCGACTTCG	CCTGCAGAGA	CAATCATTTG	ATCCA	AATCT	10560
AGAGCAATTT	TATCGATAAA	AATGGGATAA	TTGTGTTCT	GACGAATTCC	GACAGGATTA			10620
TTGGCTCCAT	GAAT	GTAACC	AGTGTTTTT	TCTAAGTCCT	TTTGTGGAAT	CATG	GCTCACT	10680
TTTTTATTGC	CAGAAATT	TTT	AGCTAGTTTC	TTTCAGACA	AGTGCTGAGT	GATA	AGGGACA	10740
ATTCCGATAAA	TCGGTCCGGT	CTTGTCTCCC	AAAAGCGCCA	AGGTTTGAA	AATCTGATCT			10800
CGTTCTAAC	CTTGAGGAAG	CTTCCTTCT	AGGGCATTGA	TTTGAATCCC	CTGATGAGGG			10860
ATAGCTGCTT	TAGATAGGAT	TTGTTCCACC	AATGTTTTT	TGATTTAAC	TTTTTTGCC			10920
ATTATTTATA	TTTATCCTCC	AATTGACTCA	TCCAAATACC	AAGCCAGATT	CCCAGCGCAA			10980
AGAAGAAGGC	GATGATGACA	TAACCGACAA	GTGAAAGTCC	TGTGTATTGG	ATACTTCAG			11040
CGTTTCCTGC	ATTGGAATT	AAGATCAAA	GGGTACTTGA	TAGGACGATA	CCGATGATGA			11100
AATGATAGAC	GAAC	GTGTTTA	CGGAGTTCTT	CTAGTTCTCC	GTCCGTCCAA	GC	GTAGGCCA	11160
CTTCTTCTTT	CTTGCTTTA	CCTTGAC	TCTGTAAAG	AGGTGGGAGG	GCAATATAGA			11220
CATGACCTGC	CTCGACTAGC	GGACGCATGT	AACGGTAGAA	AAATGTCAAG	AGCAAGGTCT			11280
GGATATGGGC	ACCGTCCGTA	TCCGCATCG						11309

(2) INFORMATION FOR SEQ ID NO: 109:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5548 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 109:

CCATAGTC	ACAAGTCTT	GTAAAGGTTT	ATCCCTGATT	CATGAAAGA	TTGTGTAAAG	60
AATCAAAAAA	AGCCACTTTT	GAAAAATGGC	TGCTCCTAAA	AATAGCTTTA	AAAATTATTA	120
GTCCTGTGCG	AAAGATTGGT	TAGGAAGAAA	AATCGTGAAG	CAACTGCCTC	TGCCAAGCTG	180
ACTCGTCACC	GTGACTGGC	CACCTAATAA	TTGACTGAGT	TCTTGACAA	TGGCAAGGCC	240
AAGACCAGTG	CCACCA	GTCTGCTTCG	ACCTTTATTA	ACTCGTAA	AACGTTCAAA	300
AATACGATCC	TGCTCTAATT	GACTAATACC	AATCCCTGTA	TCTGATACAG	AAATCTTAAT	360
GCCTTCGTC	ACCTTTGGG	TCTTGACCTC	AATTTTCCC	CCTTGTTCA	TGTAACGGAT	420

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GGCATTGGAT AAAAGATTGA GTAAGATTTG GGAAAGTAAT TGACTATCTG ATACGGAGGT	480
GACATCATCT GGCACCTGCA CCTTTAGCTG TAAATCCTTC TTCTTGAGCT GAGGTTGCAA	540
GCTTGAGTC AAATCCTGTA CAAATTCTGC CAAAGAAAGG GTCGTCCATT GTATAGGCAT	600
TTGTTGAGCC TTAGATAAGG TAAGAAGATG CTCAACAATA TGCTCAAGAC GCAAACCTTC	660
TTTGTAAATA ATGTCTAGAA AGTCATCCTT GAGCGCTCT TCTTCAGCTG ACATCCCCTT	720
AATGGTTTCA GCAAAGCCCT TAATCGAAGT AACTGGTGTCT CTCATTCTAT GGGAGGCATT	780
TGAGACAAAG GCTAAATTAA ACTTTTCATA AGTTCTAACATC GTTGTAAAT CATATAGCAA	840
GACGAGCACCA GCTTCCACAG ATTGGGTGGG GCTAAAAACG GGAACGTGTG TCACCTCTAA	900
AATCAAGTCA CCCTCATGAA ACCCACTTAC TTCTTGTGTT AACCTTGTGTT TTTGATCAAA	960
GGCTTGGTGA ACTAAATTCC GAATATCCAT CCGTTGAGG TCATCAAGTG AACTTATGTC	1020
GCGTCCACA TCGGGAAAAT AATGAGGCAG AGAGCGACTG GATAATAACA TCTGACCTTG	1080
AGCGGAAACT AAAAACGTCC CCATGGTTAG GTGCGACAGA AGAACCTCCA TTGTTTCGGC	1140
TAGATCCTG TATTGCTGAT CCTGTTGGGA GACTTTGGTT TTTAGGCCAG ACACATACTG	1200
AGCCAAAGAC TTTAAGTCTT CTTGCCCTTT TTCTAAAAAG TATTCACTAC TGGTCAAGAG	1260
AGGTTGGTGC AAGGTCTCAA AAGCAACTTC CCATTTCCAA AGGCAAAAGA GCCAGTAGCC	1320
ACCTAGTCCC AAAGAAAGGG CTAGAAGAAA GAGACCGATG CCTTTACTGA TCCAAGTTAA	1380
TGCCATCCCT GCAATCAGAA TGAGGCTAAC ACTTAGATTG ACTAGCCAAA ATTGAAGGTA	1440
GCGTTTCATC TATAACTCCT TGAACTTATA ACCATAACCC CGAATGGTTC GAATAAATTG	1500
AGGGGTTTA GGATTGTCTT CAATTTTTTC CCTCAACTTA CCAATATGAA CGTCCACCAA	1560
ACGTGTTCC TGCCCCAAAGT CATAACCCCA GATACGTTCC AAAAGACGCT CTCTAGTCAG	1620
TGTCATGTTG GGATGTTCA TAAGATAGAG CAAGAGTTCA AATTCTTTG GGGTCAAAC	1680
CAGTAACCTTA TTCGCCTTGT AGACTTCATG ACGCTCAGGG TATACTTCA AGGTCCCAA	1740
TAGCCAAGAA TCGTCAGCGA TATTATCTGA ATCATCTCCT TCTTGTCTC CTTAGTTCG	1800
CCTGAGGACA GCCTTGACAC GCGCCAGCAA TTCTCTAGGG CTAAAAGGCT TGGTCAGGTA	1860
GTCATCAGCC CCTAATTCCA AGGCCAAAC CTTATCAAAT TCATCACTTT TCGCAGAAC	1920
CATCATAATT GGAGTTTGA CGCCTTGGC TCTCAGCCGC TTACAAACTT CCATGCCATC	1980
TAATTGTGGT AACATGATAT CAAGCAAGAT AAAATCAAAG GGTTCTGTTT CTGCCAAAGC	2040
TAAGGCCTTC CGTCCATTG TCACCAATTG AGTAGAAAAG CCTTCCTTAC TTAAATGGTA	2100
GTCAAGCAAT TTCAGAATGT GTTCTTCATC ATCCACTAAT AAGACTTGTT TTGTCATCTA	2160

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TTATCTCCTA TTGGTAACAT TATAACACAA TTATCAGAAA TCCTAACATT GCTAAATCAG	2220
ATTAATTTG CCTATCAAGA CTAGTATCTG GTCAAACGCT CAATCATCTC CTTGTGCTCT	2280
GGATAGGTCG CCAGTAGATC TACCCCTTCA AATAATTCAA AATCCTCAA TTCAAAACCA	2340
GGAGCAACAA GACAAGAAC CAGAGCATCA TCCTTATCAA CTGTTGATCC CCAAATAGTG	2400
CCCTTAGGAA CACAGTAGTG AAGTTGTTGC CCTTTGGATA TGTCCAGGCC TAAAGTGACT	2460
GCTTCGTAGT GACCACATCTGC TGTAATCATG TGAACAGTAA GTGGGGATCC TGCAATGAAAA	2520
TACCAAGATT CATCTGCTGT CAATCGGTGA AAATGTGAAG GATTCGTTTC TTCTAATAAG	2580
AAATAAAATAC TGGTATAAAG CGCCCTTCCC TTACCAGCAA GGTTTATAGT GTCTGAAGCT	2640
TTTTTTGTTT GTCTAAAATA GCCACCTTCA ATATGGGGAG CTAACTCTAG AGTTCTTATC	2700
AAGTCTCTT TATCCGTCGG AGCCAATGGG TTGAAGTAAC TCTTGTCAA AGTGGTTTA	2760
CGATTTCAAG AACTCCCTCTC AGTTCTGAGG ACACGGTAAT GATTGATGCG ACGGAAGTAC	2820
AAATCAATCG CCCTAAAAAA AGAATTAGCG AATGATTCTG GTAAAAAAAAA TGCCACGCTA	2880
TGAAGGCTCA AGCGATTGTC ACAAGTCAAG GGAGAATTGT TTCTTGGAT ATCGCTGTGA	2940
ACTATTGTCA TGATATGAAG TTGTTCAAAA TGAGTCGAG AAATATCGGA CAAGCTGGTA	3000
AAATCTTGGC TGACAGTGGT TATCAAGGGC TCATGAAGAT ATATCCTCAA GCACAAACTC	3060
CACGTAAATC CAGCAAACTC AAGCCACTAA CAGTTGAAGA TAAAGCCTAT AACCATGCGC	3120
TATCCAAGGA GAGAACGAAAG GTTGGAGAAC TCTTTGCCAA AGTAAAAACG TTTAAAATGA	3180
TTTCAACAAAC CTATCGAAAT CATCGTAAAC ACTTCGGATT ACGAATGAAT TTGATTGCTG	3240
GCATTATCAA TCATGAACTA GGATTCTAGT TTTGCAGGAA GTCTATTATT TGGTTAGGTG	3300
AATTAGTGAA GCGTTTAGGC AAGTGTCTCT GGTTACGAGC TCATGGACTC TAAATCGATT	3360
ATATTTAGGG GTCATGACTA GTGAAGCAGT TAGCTAGTTC GCATATAAGC GGCTAGCGTC	3420
TAACAAATTAG GAACTTTAGT TCCAATAACT TTAAGATTAC GACGTTTTAG GACATAAAC	3480
GATCATATTT ATGTCCTAAA ACTAGTGAAG CGCCTAGCCA AAGTCCGAAT AGGATTTGGC	3540
GTTAGTTACT TAGATTGCTT TGCAATCAAG TAACTTGGC GATTTACATC TTCTCTGGCG	3600
CTTCTACTCC AAGCAAGCGA AGGGCTTCTT TGAGAACGAC TGCGGTTGCG TAGCTGAGGG	3660
CTAGACGGCT GTCGCGTTCT GGGCTTCAT CCAAGATACG TGTATGTGCA TAGTATTG	3720
TAAAGGATTG AGCCAGGCTA ATTGCAAATT TAGCAATGAT AGAAGGTTCA AAGTTATCTG	3780
CCGCACGGTT GATAATACGT GGGAAAGTCTT GAATGAGTTT AATGATTTC CAGCTTTCAG	3840
TATCATTCAA GCTATAGTTG CCAGCTGTTT CTGGTTTGAA ATCGGCTTGC CGTAAGATAG	3900
ATTGGATACG AGCGTAGGCA TATTGAACGT AAGGTCCAGT TTCACCCCTCG AAGGATACCA	3960

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TAGCCTCTAG	GTCGAAGTCG	TATCCATTG	TACGGTCGGT	TTTGAGGTCA	TAGAATTAA	4020
TGGCTCAAAT	CCCAACAGCA	TGTGCTACTT	GGTCTTTGTT	TTCTAGTTCA	GGATTTTAG	4080
CCTCGATTG	GACCTTGGCA	CGGCTAACAG	CCTCTGCAAC	AGTAGGGCTCT	AGCAAGATGA	4140
CATTCCCTT	ACGAGTAGAG	AGTTTCTTCC	CTTCTTTGT	AACCAAACCA	AAAGGAACGT	4200
GAGTAATGTC	GTCACTCCAG	TCGGTAGCCCC	TCTCTTGCAA	GACAGCTTG	AGCTGTTAA	4260
AGTGGGCAGA	TTGTTCTTGA	CCAACGACAT	AGATAGATT	AGCAAATTGG	TATTCGTTTT	4320
TACGGTAGAG	GGCTGCAGCC	AAGTCACGTG	TGATATAGAG	AGTTGCACCA	TCAGACTTCT	4380
TGATGAGGGC	TGGATGTTCA	ATTCATATT	TCTCAAGATT	CACAACTTGG	GCACCTTCTG	4440
ATTCAAGAAG	TAGTCCTTT	TCAGAAAGAA	TGTCTACAAC	TGCATCCATC	TTATCATTTG	4500
AGAAGGCTTC	TCCGTATAG	CTGTCAAATT	CAACCTTCAA	TTCATTGTA	AGGCCGTTAA	4560
ATTCACCAA	ACTTTCATCG	CGGAACCATT	GCCAAAGAGC	GAGAGCTTCC	TCATCTCCAT	4620
TTTCAAGTT	ACGGAACCAT	TCGCGCGCTT	CTTCATCCAA	GCTAGGGTCA	TTTCAGCTT	4680
CAGCGTTGAT	GCGGACATAG	AGTTTAAGGA	GTTCATCGAT	TGGATGAGCT	TTTACAGCTT	4740
CTTCGTCGCC	CCATTTTTG	TAGGCAACAA	TCAACATCCC	AAATTGTTA	CCCCAGTC	4800
CCAAATGGTT	GACCTTGACC	GTTTGATAAC	CGATTTTTG	GAAAATATGT	GACAAGCTAT	4860
CTCCGATAAC	AGTTGAACGC	AGGTGGCCAA	TAGAAAATGG	TTTAGGGATA	TTCGGACTAG	4920
ACATGTCGAT	AACAAACATTT	TCTGTCTTAC	CAATATTTG	GTCAGCATAG	TGTTCTTTT	4980
CA GTGGTAAAC	AGCTTGCAAT	ACTTGAGCAG	AAATGGCAGA	TTTATCAAGG	AAAAAGTTAA	5040
CGTAAGGTCC	TGTTGGCACA	ACTTTTCAA	AGGCTTGGCT	GTTCATTGTT	TCAGCCAGTT	5100
CAGCCGCAAT	CATTTGTGGT	GCTTTACGTT	CGACTTTGC	AAGAGAAAAA	GCAGGGAAAG	5160
CAATGTCCTCC	CATTTCTGAG	TTTTTAGGGG	TTTCCAGTAA	CTTTAAAATA	GCCTCTTGGT	5220
CCAGGCTATC	AATGATGCTA	GATAATTGCG	TAGCAATCAA	TTCTTTGTA	TTCATTAAGA	5280
GCTCCTTTT	GGACTTTCT	ACTATTTTAT	CACAATTAA	AAGAAAGAAG	AAAAAATTAA	5340
TGAAATCTCC	TGTTTTTTG	GTATAATATG	GTTATAAATA	TAGTTATAAA	TATGCACGCA	5400
AGAGGATTTT	ATGAGAAAAA	GAGATCGTCA	TCAGTTAATA	AAAAAAATGA	TTACTGAGGA	5460
GAAATTAAGT	ACACAAAAAG	AAATTCAAGA	TCGGTTGGAG	GCGCACAATG	TTTGTTGTGAC	5520
GCAGACAAACC	TTGTCCTCGTG	ATTGCGG				5548

(2) INFORMATION FOR SEQ ID NO: 110:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 3132 base pairs

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(B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 110:

TACCCGGTAG	TCTTAGCAGA	CACATCTAGC	TCTGAAGATG	CTTTAACAT	CTCTGATAAA	60
GAAAAAGTAG	CAGAAAATAA	AGAGAACAT	GAAAATATCC	ATAGTGCTAT	GGAAACTTCA	120
CAGGATTTTA	AAGAGAAGAA	AACAGCAGTC	ATTAAGGAAA	AAGAAGTTGT	TAGTAAAAAT	180
CCTGTGATAG	ACAATAACAC	TAGCAATGAA	GAAGCAAAAA	TCAAAGAAGA	AAATTCCAAT	240
AAATCCCAAG	GAGATTATAC	GGACTCATTT	GTGAATAAAA	ACACAGAAAA	TCCCCAAAAAA	300
GAAGATAAAAG	TTGTCTATAT	TGCTGAATTT	AAAGATAAAAG	AATCTGGAGA	AAAAGCAATC	360
AAGGAACTAT	CCAGCTTAA	GAATACAAAA	GTTTATATA	CTTATGATAG	AATTTTTAAC	420
GGTAGTGCCA	TAGAAACAAAC	TCCAGATAAC	TTGGACAAAA	TTAAACAAAT	AGAAGGTATT	480
TCATCGGTG	AAAGGCACA	AAAAGTCCAA	CCCATGATGA	ATCATGCCAG	AAAGGAAATT	540
GGACTTGAGG	AAGCTATTGA	TTACCTAAAG	TCTATCAATG	CTCCGTTGG	GAAAAATT	600
GATGGTAGAG	GTATGGTCAT	TTCAAATATC	GATACTGGAA	CAGATTATAG	ACATAAGGCT	660
ATGAGAATCG	ATGATGATGC	CAAAGCCTCA	ATGAGATTAA	AAAAAGAAGA	CTTAAAAGGC	720
ACTGATAAAA	ATTATTGGTT	GAGTGATAAA	ATCCCTCATG	CGTTCAATTA	TTATAATGGT	780
GGCAAAATCA	CTGTAGAAAA	ATATGATGAT	GGAAGGGATT	ATTTTGACCC	ACATGGGATG	840
CATATTGCAG	GGATTCTTGC	TGGAAATGAT	ACTGAACAAG	ACATCAAAAA	CTTTAACGGC	900
ATAGATGGAA	TTGCACCTAA	TGCACAAATT	TTCTCTTACA	AAATGTATTC	TGACCGCAGGA	960
TCTGGGTTTG	CGGGTGATGA	AACAATGTTT	CATGCTATTG	AAGATTCTAT	CAAACACAAAC	1020
GTTGATGTTG	TTTCGGTATC	ATCTGGTTTT	ACAGGAACAG	GTCTTGTAGG	TGAGAAATAT	1080
TGGCAAGCTA	TTCGGGCATT	AAGAAAAGCA	GGCATTCCAA	TGGTTGTCGC	TACGGGTAA	1140
TATGCGACTT	CTGCTTCAAG	TTCTTCATGG	GATTTAGTAG	CAAATAATCA	TCTGAAAATG	1200
ACCGACACTG	GAATGTAAAC	ACGAACATGCA	GCACATGAAG	ATGCGATAGC	GGTCGCTTCT	1260
GCTAAAAATC	AAACAGTTGA	GTTTGATAAA	GTAAACATAG	GTGGAGAAAG	TTTTAAATAC	1320
AGAAATATAG	GGGCCTTTT	CGATAAGAGT	AAAATCACAA	CAAATGAAGA	TGGAACAAAA	1380
GCTCCTAGTA	AATTAAAATT	TGTATATATA	GGCAAGGGGC	AAGACCAAGA	TTTGATAGGT	1440
TTGGATCTTA	GGGGCAAAAT	TGCAAGTAATG	GATAGAATT	ATACAAAGGA	TTTAAAAAAT	1500
GCTTTAAAAA	AAGCTATGGA	TAAGGGTGCA	CGCGCCATTA	TGGTTGTAAA	TACTGTAAAT	1560

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TACTACAATA GAGATAATTG GACAGAGCTT CCAGCTATGG GATATGAAGC GGATGAAGGT	1620
ACTAAAAGTC AAGTGTTC AATTCAGGA GATGATGGTG TAAAGCTATG GAACATGATT	1680
AATCCTGATA AAAAAGTGA AGTCAAAAGA AATAATAAAG AAGATTTAA AGATAAAATTG	1740
GAGCAATACT ATCCAATTGA TATGGAAAGT TTTAATTCCA ACAAAACCGAA TGTAGGTGAC	1800
GAAAAAGAGA TTGACTTTAA GTTGCACCT GACACAGACA AAGAACTCTA TAAAGAAGAT	1860
ATCATCGTTC CAGCAGGATC TACATCTTGG GGGCCAAGAA TAGATTACT TTTAAAACCC	1920
GATGTTTCAG CACCTGGTAA AAATATTAAA TCCACGCTTA ATGTTATTAA TGGCAAATCA	1980
ACTTATGGCT ATATGTCAGG AACTAGTATG GCGACTCCAA TCGTGGCAGC TTCTACTGTT	2040
TTGATTAGAC CGAAATTAAA GGAAATGCTT GAAAGACCTG TATTGAAAAA TCTTAAGGGA	2100
GATGACAAAA TAGATCTTAC AAGTCTTACA AAAATTGCC C TACAAAATAC TGCGCGACCT	2160
ATGATGGATG CAACTCTTGT GAAAGAAAAA AGTCAATACT TTGCATCAC C TAGACAACAG	2220
GGAGCAGGCC TAATTAATGT GCCCAATGCT TTGAGAAATG AAGTTGTAGC AACTTTCAAA	2280
AACACTGATT CTAAAGGTTT GGTAAACTCA TATGGTCCA TTTCTCTTAA AGAAATAAAA	2340
GGTGATAAAA AATACTTTAC AATCAAGCTT CACAATACAT CAAACAGACC TTTGACTTTT	2400
AAAGTTTCAG CATCAGCGAT AACTACAGAT TCTCTAAC TG ACAGATTTAA ACTTGATGAA	2460
ACATATAAAAG ATGAAAATC TCCAGATGGT AAGCAAATTG TTCCAGAAAT TCACCCAGAA	2520
AAAGTCAAAG GAGCAAATAT CACATTTGAG CATGATACTT TCACATAGG CGCAAATTCT	2580
AGCTTTGATT TGAATGCGGT TATAATGTT GGAGAGGCCA AAAACAAAAA TAAATTTGTA	2640
GAATCATTTA TTCATTTGA GTCAGTGGAA GCGATGGAAG CTCTAAACTC CAGGGGAAG	2700
AAAATAAACT TCCAACCTTC TTTGTCGATG CCTCTAACTG GATTTGCTGG GAATTGGAAC	2760
CACGAACCAA TCCTTGATAA ATGGGCTTGG GAAGAAGGGT CAAGATCAA AACACTGGGA	2820
GGTTATGATG ATGATGGTAA ACCGAAAATT CCAGGAACCT TAAATAAGGG AATTGGTGGAA	2880
GAACATGGTA TAGATAAAATT TAATCCAGCA GGAGTTATAC AAAATAGAAA AGATAAAAAT	2940
ACAAACATCCC TGGATAAAAA TCCAGAATTA TTTGCTTCA ATAACGAAGG GATCAACGCT	3000
CCATCATCAA GTGGTTCTAA GATTGCTAAC ATTTATCCTT TAGATTCAA TGGAAATCCT	3060
CAAGATGCTC AACTGAAAG AGGATTAACA CCTTCTCCAC TTGTATTAAG AAGTGCAGAA	3120
GAAGGATTGA TT	3132

(2) INFORMATION FOR SEQ ID NO: 111:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14672 base pairs

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(B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 111:

CGAGATTCT	TTAAATGAAC	TACGTGAAAT	CTACCCATCA	TCCAGATCTG	GATATTCTCT	60
CCTATCTATA	AGTAAAGTTT	TAGGAGATTT	TAATATAAGT	TCTCATGCTT	TTAAAGCTTC	120
GGTAAGAGAT	TTAAAACCGC	TCAGTTCCC	ACTCATTTGC	TTCTGGGAGA	GTTCTCATTT	180
TATTATTCTT	AAAAAAATTA	GTAAAAACAA	GTTTTATATT	TTAGATCCTG	CAAAAGGCAG	240
GCAGAGAATG	TCAATAAGTG	AATTGAAAG	GCATTATTCA	AATATCATT	TAACATTTAA	300
AAAGTTAGAT	AGCTTTATGT	CTCGTAAAGA	TAATAAGAAG	TCGCCTGTTT	TAAAGTATT	360
TTTTAAGTAT	AGGAATAAGC	TAGGGATTTT	ATTTTTGTA	ACAGCATTAT	TGTATGTAAT	420
ACAATCATT	GTACCTATAG	CTAATAGATA	CATAATTGAC	ACGAATTCA	AGGACGATTC	480
GTATTCGTCT	AGAATGTTAT	TTACTATATT	ATTTATATT	ACTGTTTCAT	TCTCACTAAT	540
GTATTTATTA	AGACAGATAT	ATGTTGCATC	CTTAAATAT	ATAATGGATA	AAGAGATTAG	600
CTATGATTTT	ATGAAACATT	TGATATATT	ACCTTACAGT	TTTTATGAAA	AACGTACTTT	660
AGGGGATATA	CTTTTAGAG	CTAACTCTAT	TGTTTATATA	AGAGAAATAC	TATCAAATAA	720
TTTTATAGCA	GCTATACTTG	ATTTGTTAAT	GATTGTGGTT	TATGCTGTGG	TTTTATTTAG	780
CTTTTCTAACG	TACATGGTAA	TCTTTTTAAT	ATCACTAAGT	CTAGCTCTAT	CTATTGTAAT	840
GTATCCAATC	ATAAAAATCT	CAAAAAATT	AATTGATAAA	AATATAAAAG	AAAAGTTAA	900
TGTTCAAAAT	ATTACTTCCG	AAGTAATTTC	AAAAAATAGT	GATATTAAGC	TAACGGAGA	960
AGAGGAATT	TGGATTAACA	AATGGGATAA	TTTTAATACA	AAACAGCTCA	TCATAGGTG	1020
AAAACTTGAT	ATACATTAT	CAATTGTTAG	TAGTATAACG	AATGTTTAC	AAATTATTCT	1080
CCCTGTTTG	ACCCTTATTG	TAGGTGTAAA	TATAAAACA	TTCGAACAA	TGACGTTAGG	1140
ACAAATTGTA	GCAATAAGTA	CAGTCTCACC	ATACTTTATT	TCTCCTATAA	TTTCTTTAAG	1200
TGATAACTAT	ATACAATTAA	TGTTATTAAA	GGGATATT	TTAAGAATAG	AGGATGTGTT	1260
TAATACTAAA	TCCGAATTAA	TTCCAGAAAG	AGTCAGTC	GATATAAAAT	TTGATAAAAA	1320
AATAGAATTAA	AAAGATATT	GGTATAAAATA	TGGATTATT	GATGATTATG	TTTGAAAGG	1380
AATAAAATGTT	ACTATTAAAA	AAGGAGAAAC	TGTTGCTATT	GTTGGAGAAT	CAGGTTCAAG	1440
TAAGAGTACA	TTAGCTAAA	TTTATTAGG	TTTATTAGAA	CCTAATATTG	GTTCAATAGA	1500
AGTTGATGGA	GTAGAAAAAG	AAGAAATTGG	TCAACATTG	TATAGAAAGA	TTTTGGAGC	1560

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AGTGTACAA AATTCAACCC TAAGTTATGG TACCTTAAGA GAGAATTGGA CATTGGACA	1620
CTTGTTTCA GATGAAGAAC TAATGACAAA TCTAAATTCA ATTGGTCTTA GCAATGTAGT	1680
TAAATCTTA CCTCTGGAT TAGAGACAAT CATCGCTGAA GAAGGTAATA ACTTTCTGG	1740
AGGGCAGCAG CAAATGATAC TTTTAGCTCG TTGTCTTTG TCGAACCTT CGGTAGTTGT	1800
TTTGGACGAA GCAACAAGTA GTTTAGATAA TTTATCTCAA CAAATTACAA CTTCTTACTT	1860
AAGTGAATC GGTACCACTA AGATTTAAC TGCCCACGAA CTAGATACTA TCAAGTCTGC	1920
AGATAAGATC TTAGTAATGC ATAATGGTGA AATTGTAGAG ATTGGGACCC ATAGAGAACT	1980
TCTTGAACTA GGAGGCATTT ATAAGCAATT GTATTCAAAT AATTAGTTT TGATTAAG	2040
GGTAAATTGTA TGAAGATTAT GAAAAAAA TATTGGACTT TAGCGATATT ATTCTTTGT	2100
TTGTTCAATA ATTCTGTTAC TGCTCAAGAA ATACCTAAA ATCTTGATGG CAATATAACT	2160
CACACTCAGA CTAGCGAAAG TTTTCTGAA TCTGATGAAA AACAGGTTGA CTATTCTAAT	2220
AAAAATCAAG AAGAAAGTAGA CAAAATAAA TTTCGTATTC AAATCGATAA GACAGAATT	2280
TTTGTAAACAA CAGATAAACAA TTTAGAAAAA AACTGTTGTA AATTGGAAC TGAACCACAA	2340
ATAAAATAACG ATATTGTTAA CTCTGAAAGT AATAATTAC TAGGCGAAGA TAATTTAGAT	2400
AATAAAATTAA AGGAAAATGT TTCTCATCTA GATAATAGAG GAGGAAATAT AGAGCATGAC	2460
AAAGATAACT TAGAATCGTC GATTGTAAGA AAATATGAAT GGGATATAGA TAAAGTTACT	2520
GGTGGAGGCG AAAGTTATAA ATTATATTCT AAAAGTAATT CTAAAGTTTC AATTGCTATT	2580
TTAGATTTCAG GAGTCGATTT ACAAAATACT GGATTACTGA AAAATCTTC AAATCACTCA	2640
AAAAACTATG TCCCCAATAA AGGATATTAA GGAAAAGAGG AGGGAGAGGA AGGAATAATA	2700
TCAGATATTC AAGATAGATT AGGTCACTGGT ACGGCTGTTG TAGCTCAAAT TGAGGGAT	2760
GACAATATTA ATGGAGTAA TCCCTCACGTT AATATTAACG TCTATAGAAT ATTTGGTAAG	2820
TCGTCAGCTA GTCCAGATTG GATTGAAAA GCAATTGTTG ATGCTGTAGA TGATGGCAAT	2880
GATATTATCA ATCTTAGTAC TGGACAATAT TTAATGATTG ATGGAGAATA TGAGGACGGA	2940
ACAAATGATT TTGAAACATT TTTGAAAGTAT AAAAAGGCTA TTGATTACGC GAATCAAAA	3000
GGAGTAATTA TAGTAGCTGC ATTAGGGAAT GACTCCCTAA ATGTATCAA TCAGTCAGAT	3060
TTATTGAAAC TTATTAGTTC ACGCAAAAAA GTAAGAAAAC CAGGATTAGT AGTTGATGTT	3120
CCAAGTTATT TCTCATCTAC AATTCGGTC GGAGGCATAG ATCGCTTAGG TAATTATCA	3180
GATTTTAGCA ATAAAGGGGA TTCTGATGCA ATATATGCGC CTGCAGGCTC AACATTATCT	3240
CTTTCAGAAT TAGGACTTAA TAACTTATT AATGCAGAAA AATATAAAGA AGATTGGATT	3300

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TTTTCGGCAA CACTAGGAGG ATATACGTAT	CTTTATGGAA ACTCATTGCG	TGCTCCTAAA	3360
GTTTCTGGTG CGATTGCAAT GATTATTGAT	AAATACAAAT TAAAAGATCA	GCCCTATAAT	3420
TATATGTTG TAAAAAAATT CTGGAAGAAA	CATTACCAGT AAAAAATGGT	ATAAAAGTGT	3480
TAAATATACC AAACGTATTG AGATATGATT	TGAATATGTT ACAATTAGAA	TATAAAAATG	3540
AACAAAGTTG GGATAGTTTC ATAGATAATG	TTAATTAAAT TGAGTTGGAA	GAGAGAATTC	3600
AAACTACTAT TGGAATTAAA CAAATAAACAA	CACACAATAT TATTACTATT	GCCCCGAGAAG	3660
GGTACTCTCA AAATTATTTA CCTAACACTT	CAGAAAATAC ATATAATTCA	TTACAAGTCA	3720
GTTTAGTTGG AGTATTACTA CTTTTATCAA	GTATGGTAAA TATTTTATGG	GCTAAAAAAA	3780
GTAAATGAAA ATAAAATTG GAGCCCTCTG	AAAAAGTAAG TCCTACAGTT	CAACTAAAAT	3840
GAGTCAAAAG ATGAATCACCA TTGATGTAGG	GGAGTTTGTC TTATTGCTGC	CTGAACACCT	3900
CCGTTTCAGAG GAAGAACATT ATAAATCTGT	TTTGAAGAC GACTTAACCA	GTCGCATATC	3960
TAGTCAAGAT GAACGACAGC AAATGACTGC	TACGGTAGGT TATTTAGAAT	CAGGTCAGGA	4020
TCGTTTGTG TATAATACGA CCCCTATTTC	TTACCAGCAG TTTTGAAAG ATCCAATCAT	4080	
CATTGTTATA ACACCCCAAT CAACTGGTCC	ACAGTCCATT TTGTTTGGA	TAGACGCAGT	4140
ACAGAACTAC GTTCTCTTA ATCAATTGTC	TGATGCCAG GAGCTTATCC	AGAGACAAAGG	4200
CATTGAAAAT TGGGTCAG AAATGCAAAC	AGGTTACCAC AACTACATCA	CATTATTGGA	4260
TAATATCCAG AGGGAACGTT GGGTAATGCT	AGCAGGAGCT GTGCTTGGGA	TTGCAACTTC	4320
AATCTTGTG TTTAACACTA TGAATAGGCT	CTACTTTGAA	GAATTTAGAC GTGCCATT	4380
TATCAAACGC ATTGCAGGTC TCAGGTTCTT	AGAAATCCAT CGCACTTATC	TCTTGCTCA	4440
ACTGGGTGTG TTTTTACTGG GATTGTTGC	GAGTGTATTT CTTCAGGTAG	AGATAGGAGT	4500
TGCTTTCTTA GTCTTGTAC TCTTTACTGG	TCTATCTCTT TTACAGTTAC	ATGTCCAAT	4560
GCAGAAAGAA AACAAAGATGT CCATGCTTGT	TTTGAAGGGAA	GGTTAATATG ATTGAACCTTA	4620
AACAGGTGAG TAAATCTTT GGAGAACGAG	AGTTATTTTC	GAATCTTCA ATGACATTG	4680
AGGCTGGAAA AGTCTATGCC TTAATTGGTT	CAAGTGGTAG CGGAAAAAACAA	ACCTTGATGA	4740
ACATGATTGG GAAATTAGAA CCTTATGATG	GGACGATTTT TTACCGAGGT	AAAGACTTGG	4800
CCAATTATAA ATCAAGTGAT TTTTCCGTC	ACGAATTGGG CTACCTCTTC	CAGAACTTTG	4860
GCTTAATTGA AAACCAAAGT ATTGAAGAAA	ACCTTAAGCT AGGTCTCATT	GGTCAAAGT	4920
TGAGTCGGTC GGAACAGCGG TTGAGGCAGA	AGCAGGCTTT AGAACAGGTC	GGCCTGGTT	4980
ATCTTGACCT AGATAAGCGC ATCTTGAGT	TATCGGGCGG AGAATCGCAA	CGGGTTGCCT	5040
TGGCAAAAT TATCTAAAG AATCCACCC	TTATTCTGGC AGATGAGCCA	ACAGCTTCAA	5100

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TAGACCCAGC AACCTCTCAG TTGATTATGG AGATTTGCT ATCTCTTCGA GATGATAATA	5160
GGCTAATCAT TATCGCAACA CATAATCCGG CAATTTGGGA GATGGCTGAT GAAGTGTCA	5220
CGATGGATCA TCTGAAATAA AAATCCTTGT TTTTAATTGC ACGATGAGTT ACTGAAATAT	5280
TATCATGAAT CAAGAATTGG AGTTAATTAA GAATTGTACT TAATTTAGAA TTGTACTTTA	5340
TTAATATTGA GGTAACCTTT TCTTGATAAA GGAAGAAATA ATGGAGAGGA AGTTAGAATG	5400
AAAAAATTG ACAATTATAT TATTGAGAAG CCTTGCGATT CTAATTCAA TAAACTGCAA	5460
AAAATCTAA TAATTGAAAG TTTGGTAGAT GATATTTGC AATTTCTCT CAGAACATCAAT	5520
AATAGTGTAG GAGAGATTT CCTCCTACAA CCGTTTAAAG AGAAAACAT CTTTATTCCA	5580
TGTTATTTG AGGAAGATAT TGTGAAAGTC AAAGATGATG ATAAAGTTGA GTGGAATTTG	5640
TTAGAATTTC AAAAATTTAG AGCATTTTG GCTTAGTAAT CTGTGTTGAA GGCTCAAAAC	5700
CTATGGTAA AAAGTAGCTT TGAAAACGTA TTGCTCCAA AGATTTAGTT AAATAATGAT	5760
TTAACACAAA AAGAAATTAT TGAAGTTCTG GAAAGATGTT GTTTCAGTAT TGAGAAAAGG	5820
TGGGAAAAAC TTGCGATTTT CACAGAGAAA GGAAGAAAAA GTATAGAAAT ATAGTCAATT	5880
GAAACAAGAA CAGGATAAAA GAACCTTTG TGCCATATT TTCTCCTTTC GCTTACAAT	5940
TGGATTGAAC ACCTTTATTG TATCGCGTTT GGAGTTTTT TGGTATAACC TTCGACGCAC	6000
ACCCGCATAG CGGGTGTGTTT TTTTGTCTCG CACCTAACGG AGCGAGACAA ACTAATAGTC	6060
ACTTAATCAA AAAACGCACC ATATCAAAAAA CTAAAAAGTT TGATATCATG CGTCATGTCT	6120
TAAACTAATT GACTATACTT TCTATTCAA TGAGCTTTA ACCAATTGAT TGAGCCAATC	6180
CACTCTAAA ACCAAAGACC AATTCTCGC TTGACTGACT CTTCTGAATC TGAACCATGT	6240
ACAACATTTC GGATAATCTC ATTTCTCCA GCAGCTTTG CAAAATCACC TCGAATAGTG	6300
CCTGGTAAAG CTTCTCTGG ACGAGTTGCA CCCATCATGG TCCGCCAAGT TTCGATTACT	6360
TTGGGACCAAG AAATGACACC CACAAGAACT GGACCTGAAG TCATGAATTC ACGAATCGGT	6420
GGGTAAAAAC TCTGACCAAC CAAGTCCTGA TAGTGCTGGT CAATCAACTC TTCTGAAACC	6480
TGTGAACGAA ACTCCAATT TTGATTGTA AATCCACGTT GTTCGATGCG CTTAACACT	6540
TCACCCACTA GCCCTTTT TACACCCTCT GGTTTGATGA TAAAGAATGT TTGTTCCATA	6600
CCCGTCTCCT TTGTCAGCTT CTTCTTTA TTTTACCAACA TTTCGTGGAA AAATGGAGAA	6660
AGTTTCAGA AGAGAGAATG AGAGAACCT CGGGTTCTCT CATTCTCTCT TATTCTACTG	6720
TTTCTTCCAC AGTTCAACG GCAGTATCCA CAACTACTTC TGTTGTTCT TCATTCCTT	6780
CTTCCTCTAC TGGAGGATTA AGGTATTCTT CTTCGTGAC AGCATGTGGT TCAAGGTTAC	6840

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GGTAACGGGC CATAACCAGTA CCAGCTGGGA TGATCTTACC GATGATAACA TTTTCTTTAA	6900
GTCCAAGGAG ATGGCTTTTC TTACCACGGA TAGCTGCGTC AGTAAGGACA CGAGTTGTT	6960
CCTGGAAGGA AGCCGCTGAC AAGAAACTGT TTGTTCAAG TGAGGCTTTG GTAATTCCCA	7020
TAAGGACTGG GCGACCTGTC GCTGGAACCTC CACCTGCGAT AAGGACATCT TTGTTGGCAT	7080
CTGTAAAGTC ATTGATATCC ATGAGGGTAC CCATGAGAAG ATCTGTATCA CCTGGATCCA	7140
TGACACGGAC TTTACGGATC ATTGACGAA CCATTACCTC GATGTGTTG TCACCGATTT	7200
CTACCCCTTG GCTACGGTAA ACTTTTGTA CTTCACCGAG AAGGTACGTT TCAACTGACA	7260
AGACATCACG AACTGCAAGG AGACGTTTG GTTGGATAGA ACCTTCTGTC AGAGCAGCAC	7320
CACCGCGCTAC TTGGCCCCCA ACTTCGACAC GCATACGAGC TGTAATGGA ACGACATATT	7380
CACCTTCGCC AGTTTCACCC TTAACAAAGA CTTTCTTGGT ACGAGTTGAT GCATCTTCTT	7440
CGATAGCAGT AACTTGTCTT TTAACCTCTG TAATAACCGC TTCCCCTTTA GGATTGCGGG	7500
CTTCAAAGAT TTCTGGACA CGAGGAAGAC CCTGAGTGAT ATCGGTATTT GAGGCAACCC	7560
CACCTGTGTG GAAGGTACGC ATTGTAAGCT GTGTACCAGG TTCCCCGATA GATTGGGCAG	7620
CGATTGTACC AACTGCTTCA CCAACTTCAA CGCGATCACC AGTCGCCAAG TTGATACCGT	7680
AACAGTGACG GCAGACACCG TGACGAGTGT TACATGTAAA TACAGAACGG ATAGTCACCT	7740
CTTCCACACC AGCATTGACA ATTTCACGCG CCTTGTCTTC TGTAATCAAT TCATTTGGAC	7800
CAATAATCAC TGCACCAAGTT TCTGGATGTT TAACAGTTTT CTTAGTGTAA CGACCGTTGA	7860
GACGCTCTTC GAGAGACTCG ATCATCTCTT TTCCTTCTGC GATAGAACGG ATCAAGAGAC	7920
CACGGTCAGT TCCACAGTCG TCCTCACGGA TGATAACGTC TTGGGCAACG TCGACCAAAAC	7980
GACGAGTCAA GTAACCTGAG TCGGCTGTCT TAAGGGCCGT ATCGGTACATA CCTTTACGAG	8040
CACCGTGAGT TGAGAAGAAC ATTCCAATA CCGACAAACC TTCGCGGAAG TTTGAAAGGA	8100
TTGGCAATTC CATGATACGT CCATTCGGAG CAGCCATCAG ACCACGCATA CGGGCAAGCT	8160
GTGAGAAGTT TGAGATGTTA CCACGGGCTC CAGAGTCCAT CATCATAACG ATTGGGTCT	8220
TAGGATCTTG GTTAGCAATC AAGCGTTCT CAAGTTTTTC ACGGGCAGCA CGCCATTAG	8280
CTGTAACAGC ATTGTAACGC TCGTCGTCTG TGATCATACC ACGACGGAAT TGTTGGTGA	8340
TTTGTTCGAC ACGTTGTGT GATTCTCAA TGATTCAGC CTTGTACATCA ACGACTGGGA	8400
TATCGGCAAT ACCCACTGTC AATCCTGCAA GAGTTGAGTG GTGGTAACCG AGGTTCTTCA	8460
TGCGGTCAAAG TAGGGCAGAA GTTCTGTGCG TACGGAAACG TTTGAAGATT TCAGCGATGA	8520
TATTTCCAAG GTTTCTTCTTC TTGAATGGAG GGTTGAGCTC AAGATTGCTG ATAGCTTCCT	8580
TGATATCTCC ACCAAGTGGC AAGAAGTATT TAGCTGGAAC ACCTTCTGTC AAGTTGGCAT	8640

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TGTTTGGTTC TTGCAAGTAT GGTAGCCCT CTGGCATGAT ATCGTTGAAG AGAATTTCAC	8700
CAACTGTTGT AAGCAAGACC TTATGTCCTT GCTCTCTGT CCAAGGCTTG TTGAGGCTGT	8760
CTGTTGCGAT ACCAACACGT GAGTGGAGGT GAACATAACC ATTGCGGTAA GCCATAACCG	8820
CTTCGTCACG GTCTTGAAG ACCATTCCCTT CACCTTCGCG ACCAGCTCT TCCATGGTCA	8880
AGTAGTAGTT ACCAAAACC ATGTCCTGAG ATGGAGTAAC TACCGGTTTC CCATCTTCG	8940
GGTTCAAGAT GTGCTCAGCA GCTAGCATGA GGATACGAGC TTCTGCTTGT GCTTCTCTG	9000
AAAGTGGTAC GTGGATGGCC ATTTGGTCCC CGTCAAAGTC AGCATTGTAG GCTTCACAGA	9060
CAAGTGGGTG CAAGCGAAGA GCCTTACCAT CAATCAAGAC TGGCTCGAAG GCTTGGATAC	9120
CCAAACGGTG AAGGGTCGGT GCGCGGTTCA AAAGCACTGG GTGTTCTTAAATCACCTCTT	9180
CAAGGATATC CCAGATAACGC TCATCTCCGC GTTCCACCAA GCGTTAGCT GCTTGCACGT	9240
TTTGCACGAT ATCACGGGCA ACGATTTAC GCATGACAAA TGGTTTAAAG AGTTCAATCG	9300
CCATTTCACG CGGCACACCA CATTGGTACA TCTTAAGAGT TGGACCAACG GCGATAACTG	9360
AACGTCCCTGA GAAGTCAACA CGTTTACCGA GCAAGTTTG ACGGAAGCGT CCTTGTCTTAC	9420
CTTTAAGCAT GTGGCTCAAT GATTCAATG GACGGCTACC TGTCCTGTG ATTGGACGAC	9480
CACGACGACC ATTGTCAATC AAAGCGTCAA CTGCTTCTTG AAGCATAACGC TTCTCATTTC	9540
GAACGATGAT ACCTGGTGCA TTTAACTCAA GCAACCGAGC CAAACGGTTG TTACGGTTGA	9600
TAACACGGCG GTAAAGGTCA TTCAAGTCAG ATGAGGCCAA ACGGCCACCA TCCAAGTGCA	9660
ACATTGGACG AAGATCTGGT GGGATAACCG GAAGGATGTT AAGAATCATC CATTCAAGGTT	9720
TGTTTCCAGA CTTGTAAAAG GCATCCAAAA CATCCAAACG ACGGATGGCT TTGACACGCT	9780
TTTGTCCAGT AGCTGTTTTC AATTCTTCTT TGAGTTTCAGC AATTCTTTT TCAAGATCTA	9840
CTTGCTTCAA AAGGTCTTGG ATGGCTCCG CACCCATCTT GGCAACAAAT GAACCATAAC	9900
CATATTACG CAAGCGCTCT CGGTATTCGC GCTCTGTAT GATAGACTTG TGCTCAAGTG	9960
GTGTATCCTT AGGATCAATC ACCACATAAG CCGCAAAGTA GATAACTTCC TCGAGGGCAC	10020
GAGGGCTCAT ATCAAGGGTC AAGCCCACAC GGCTTGAAT CCCCTTGAAG TACAGATGT	10080
GAGATACAGG AGCTTCAAT TCGATATGTC CCATACGCTC ACGACGAAC TTCGTACGCG	10140
TTACTTCAAC CCCACAGCGG TCACAAACAA TTCCTCTGTA ACGAATGCGT TTGTACTTAC	10200
CACAAGCACA TTCCCAGTCT TTTGTAGGAC CAAAGATCAC TTCATCAAAG AGTCCTTCAC	10260
GTTCTGGTTT CAAGGTACGA TAATTGATTG TTTCAGGTTT TTTGACTTCT CCATAAGACC	10320
ATGAACGGAC TTTACTTGGA GAAGCTAGGG TGATTTGCAT ACTTTAAAA CGATTACAT	10380

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CAACCACAT	TTCTTCCCTT	TCTATTCTAA	GTGAACTGCT	TATTCTGTGTT	CAGCAGCTTC	10440
TTCTGTTGCT	TCCGCTTTTG	TTGCTTTCTC	AGCTTCTTCA	GCTTCAAAGG	CTGCTTTAGC	10500
CTCTTGGGCT	GCTTTTCGC	GGGCTTTTC	AAGGTCATCT	ACGTGGATGA	CATCTTCGTC	10560
CATTCTTCA	TCCAAGTCGC	GAAGTTCCAC	TTCTTGGTCA	TCTTCGTCTA	GGACACGCAT	10620
GTCAAGACCA	AGAGATTGCA	ATTCTTGAC	AAGAACTCGG	AAGGATTCTG	GAACACCTGG	10680
TTTTGGAATT	GGTTTGCCTT	TTGTAATAGC	TTCATAGGCT	TTCAAACGTC	CGTTGATATC	10740
GTCCGACTTG	TAAGTCAAGA	TTTCTTGAAG	GACATTTGAC	GCACCGTAGG	CTTCAAGAGC	10800
CCAAACCTCC	ATCTCACCGA	AACGTTGTCC	ACCAAACCTGA	GCCTTACCTC	CGAGTGGTTG	10860
TTGGGTAACA	GTTGAGTATG	GTCCGACTGA	ACGCGCGTGC	AATTATCAT	CAACCATGTG	10920
GTGGAGTTG	ATCATGTACA	TGACTCCGAC	AGAAAACACGG	TTATCAAACG	GTTCACCAGT	10980
ACGTCCATCG	TAAAGGATCG	TTTTGGCATC	GCTATCCATA	CCTGTTCTT	TAACAGTTGA	11040
CCAAAGATCT	TCAGAACTTG	CTCCATCAAA	GAETGGTGT	GCGATGTGAA	TACCAAGAGT	11100
ACGAGCTGCC	ATACCAAGGT	GAAGCTCCAT	AACCTGACCG	ATATTCAAC	GTGATGGTAC	11160
CCCAAGTGGG	TTCAACATGA	TGTCGACTGG	AGTTCCGTCT	GGAAGGTAAG	GCATGTCTTC	11220
TACAGGAACG	ATACGAGAGA	CAACCCCTTT	GTTCGGTGA	CGTCCGGCCA	TTTTATCTCC	11280
GACCTTAATC	TTACGTTTT	GAGCGATGTA	AACACGAACC	AACATGTTAA	CACCTGATTG	11340
CAAATCCT	CCATTTACAC	GTGTAAAGAT	CTTAACATCA	CGAACGACAC	CATCGGCACC	11400
GTGTGGTACA	CGAAGAGAAAG	TATCACGCAC	TTCACGAGAC	TTGTCCTCAA	AGATAGCGTG	11460
CAAGAGACGT	TCTTCAGCTG	AAAGATCTTT	CTCACCCCTTA	GGTGTACTT	TACCTACAAG	11520
AATATCACCT	TCTTTAACCT	CAGCACCAAT	ACGGATAATC	CCCATTCGCT	CAAGGTCTTT	11580
GAGGGCATCT	TCACCAACGT	TTGGAATTTC	GCGAGTGATT	TCTTCAGGCC	CAAGCTTGT	11640
ATCGCGCGTT	TCTGATTCTG	ATTCTTCAAG	GTGAAACAGAT	GTGTAGACAT	CGTCCTTCAC	11700
CAAGCGTTCG	CTCATGATAA	CGGCATCCTC	GAAGTTGTA	CCTTCCCAAG	TCATGTAGGC	11760
AACGATTGGG	TTTTGTCCAA	GCGCCATTTC	TCCATTTCC	ATAGAAGGTC	CGTCAGCGAT	11820
GAAATCGCCT	TTTTCAACGA	CATCACCAAC	TTTACGAGA	GTGCGTTGGT	TGTAAGCAGT	11880
ACCTGAGTTT	GAACGACGGA	ATTTTGGAT	GTGGTAAACA	TCCAATGAAC	CATCTTCACG	11940
ACGAACCTCT	ACCTGTCAG	CATCTCGCTA	AGTAACCTTA	CCATCATACT	GAGCAATCAC	12000
AGCCGCACCA	GAATCGTGGG	CTGCTTGGTA	TTCCATACCA	GTACCAACGT	AAGGTGCCTG	12060
AGGATTAATC	AATGGCACAG	CCTGACGTTG	CATATTGGCT	CCCATGAGGG	CACGGTTGGA	12120
GTCATCGTT	TCCAAGAAAG	GAATACATGC	TGTCGCAACG	GCAACTACCT	GTGTTGGTGA	12180

813

AACGTCCATG TAGTCAACAA TATTAGCTGG ATACTCTGG TTGACCCCTT GGTGACGTCC	12240
CATGACAATC TTCTCAGCAA AGGTTCCATC TTCATTAGA CGAGAGTTAG CCTGAGCTAC	12300
AGTATATTCA TCTTCATCAGCTGTCAA CCAAACAATT TCGTTCGTGA CAACACCTGT	12360
TTCACGGTCA ACCTTACGGT ATGGTGTGG AACAAAACCA TATTGTTCA AGTGTCCATA	12420
AGATGACAAG TTATTGATCA AACCGATGTT AGGTCCCTCA GGTGTCTCGA TTGGACACAT	12480
ACGACCATAG TGAGTGTAGT GCACGTCACG TACTTCATAT CCAGCACGGT CACGAGTCAA	12540
ACCACCAGGT CCTAAGGCTG ACAAACGGCG TTTGTGAGAC AACTCAGAAA GCGGGTTGTG	12600
TTGGTCCATG AACTGTGACA ACTGTGATGA ACCAAAGAAAT TCTTTAACTG CAGCTGTTAC	12660
AGGACGGATA TTGATAATTT GTTGTGGTGT CAAGACTTCA TTGTCCTGAA CAGACATACG	12720
TTCACGGACA TTACGTTCCA TACGAGAAAG TCCCAAACGT ACTTGGTTGG CAAGCAATT	12780
ACCAACCGCA CGGATACGAC GATTCCAAG GTGGTCGATA TCATCTACAC GGCAAGTCC	12840
TTCAGCCAAG TTGAGGAAGT AGCTCATCTC AGCAAGGATA TCTGCAGGAG TCACCGTACG	12900
AACCTTGTCA TCTGGGTTAG CATTACCAAT GATCGTTACG ACGCGATCTG GATCAGTTGG	12960
AGCAATAAAC TTGAATTTT GAAGAACAAAC AGGCTCAGTC ACAACGGCTG CATCGTTGG	13020
GATGTAGACA ATCTTGTCA AGTCGCCATC CAAATGGCTT TCAATGCTTT CAATCACGCT	13080
ACGAGTCATA ATCGTACCAAG CTTCTACCAA GATTCTCCA GTTTCAGGGT CTACCAATGG	13140
CTCTGCAATG GTTGGTTGA GCAAACGTGT TTTAACATTG AGTTTTTAT TGATTTGTA	13200
ACGACCAACT GCTGCCAAGT CATAACGACG TGGGTCAAAG AAGCGAGCTA CAAGCAAGCT	13260
ACGTGAGCTT TCAGCCGTCT TAGGCTCAC TGGACGAAGG CGTTCGTAAGG TTTCTTCAA	13320
GGCTTCGTCT GTACGAGAGT CCATTGGATT CTTGTGGATA TCTTTTCAA CAGTGTGCG	13380
AAACCAATTG CTGTACCAAA AGATATCAA GATTCATCA TCACCTGAGA AACCAAGAGC	13440
ACGAACCAAG GTTGTAATG GAATCTTACG AGTACGGTCG ATACGAGTGT AGGTGATATC	13500
TTTTGAGTCG CTTCAAGTT CCAACCAAGC TCCACGGTTA GGGATAACAG TTGAACCATA	13560
GCCCACCTTA CCATTGGTCT CTACTTGTGTC GTTAAAGTAA ACACCTGGTG AGCGGACCAA	13620
CTGAGAAACG ATAATACGTT CACCACCATT GATGATGAAA GTACCCATT CTGTCATGAT	13680
TGGGAAATCA CCAAAGAAAA CTTCTGGGT CTTGATTCG CTTGTTCTT TATTGATCAA	13740
ACGGAAGGTT ACAAAAATTG GTGCTGAGTA GCTAGCATCG TGGATACGAG CTTCTTCTAG	13800
CGTATATTAA GGTCCTTGA TTTCATATCC AACAAATTCC AACTCCATTG TGTCTGTGAA	13860
GTGGAAATT GGCAATACAT CTTCAAACAC TTCCCTAAGA CCGTGGTCTA GGAAAGCTTT	13920

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GAATGAGTCA	GTTTGAAATT	CAATCAAATT	TGGTAAGTCA	AGAACTTCTT	TGATTCTTGA	13980
AAAAC TACGA	CGGGTACGAT	GTTTCCC GTA	TTGAACGTCA	TGTCCTGCCA	AGATGATTCT	14040
CCTTTGTAAA	TAAGTTCCAA	GCCTTGTCAA	TCAGGCTTTT	CTAACCGTCA	TATGGTTGTA	14100
AACCCCTTAT	CACCGTGTCC	TCTTGACGAA	TTTCAGAAT	CTTTAACGCT	CTGTTACAAA	14160
TGCTCAAAAT	CTTGAAAAAA	AGCACAAAAA	GAGCAGCTAA	ATCTGACTTT	TTCAGAAGAT	14220
TTAAC TGCTG	TGAGCCTTGT	CTGGACAATA	TTTCAGACAA	AACCTACGAC	AAATGATTAC	14280
CCATATTATA	CCCTATTTAG	CTAGATTTT	CAAGGGTTT	CAGTAGGTTT	TTGGTAAATT	14340
TTTTCCCATA	AAAAACTTGG	CATCACATTC	GAATCACGCT	ATGGTACAAA	AAACTGAAAA	14400
AACTATTGAC	TGAAAATCAT	TTTCAAGGTA	TAATAATAAA	CGTTAACGCG	GTATAGCCAA	14460
GTGGTAAGGC	ACGGCTCTGC	AAAAGCTTGA	TCGTCGGTT	AAATCCGTCT	ACCGCCTTCT	14520
ATAACTTGAT	TTATCAGGTT	TCAAATGAAC	AGAAAGCCCA	ATTTGAAGGG	CTTTTTTAT	14580
TTTCCCTCGA	ATAAAATACGT	ATAACTTTAA	AAACTTTGG	AGCGAGTTG	TGGCAGAGTT	14640
CTTCCCATGG	CATAATTCCC	TTTGAAATC	AG			14672

(2) INFORMATION FOR SEQ ID NO: 112:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7902 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 112:

AGGAGACTAT	TCAAGCCAA	ATTgAGTAC	CCAGCAAAGA	CTGTATAGAC	TGTGATACGT	60
TTTTCATAGC	CATTGGTAAA	GAGAATTG	GAACCAAGAA	TGGTATCTAA	GGCCAGGATA	120
ATCGTACGAA	AAGCGAAGAG	AGAGGTCAAG	ATGCCGCCTC	CGATATATTT	TTCACTACCG	180
TAAAGTAGGA	TGGCATTG	TCCTAAAACC	ATGAGTCAA	AACTCAGTGG	AATGATAAAG	240
AAGTTAAAGA	TTCGACTACC	TCTATTAACC	AGAGAAACAT	AGGTTCTTT	GTCTCCTT	300
CCCAGATAGT	AACTGAGACG	AGGCACACTC	ACTCCAATTG	CACCTGTTAC	AAACCCAGCT	360
ATAACGGTCA	CAATTGCTG	AGCTATGGTA	TAGTAACAA	CGTTGACATC	AATCCCTGTT	420
TTAACGAGGA	AGAGGCAGTC	AAAAAAAGTG	AAGAGCATAT	TGGCATTGGC	AAAGACTAAC	480
ATGGCTGTCA	GAGGGAGAAA	GAGTGGTTA	AAATCACTTA	GGTGAATT	AAACAGTTG	540
ATGTCTCTT	TAATCCAAA	ATAACTAATC	AGGTAGTTAA	TCAGCGTCGA	TAAACTCATC	600
ACAAGTGTAT	AGACAACAAAT	ATCGTGTCA	TTTTAACAA	ATAAGAAAAT	AGAGACCAGC	660

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ATCAGGATAAC GGATGAAGGC AGTTTGTA AAGAGAAAAC TGTAATTTC CAGAGCTCA	720
TTGACCCATT CGATTGAAAA AATCTGGCA ATGAGTTGAA TCCCCATAAC AAGGTAGACC	780
TTTTGACGA TTGGATTATC AGTAAAGAAG AGAGGATAGG CTAGGATATA GACAGCAGTG	840
GTCAAAATCG TACAAGCGAT GCACAAATAA AAAAGACTAG AAAAGGTTCT GTTAAGATCT	900
TTTTGTTAT CCTTGACATT ACTGATAGCC CTTAAACCGT AGTTATAGAC ACCATAAGTT	960
GCAAAGGGCA AGAAAAATGA CAAAATAGT TCGACTGAGT TGAAGTAACC ATAGTCAGTT	1020
CGGTCCAAGA CACGCGCAG ATAGGTTCCA GTTAGGATGG GAAAATAAT ATTCAAGACA	1080
CGAATTCCA TGTAAGATAG AGCATTAAAT TTTATACTTT TCATTCAATT TACCTCGTT	1140
TTCATTATAT CATAAAGTTA GCTAATAAGA AATGAAGGGC AGTAAGTCAA GTAATCACTT	1200
TGAAGTTCA AATCTTAAGT TTTAAGTTTT CTTTAAGGAA AGTATATTAT TCTGAAGGAC	1260
TCTAAAATTT CGCACCCATT TATTAGTAAT TGCTACAGAA TTCCTAGTCA TTACTAGAAA	1320
TGGACTAGTT TCTTGAAATA ATAGAACTGC ATAATTCTCC TATTCTAGAA GGGGAGGACC	1380
AGTATTTCTT TTATGATAGG ACTAGATTGT GGTATAATAG AGAGAATAAG TTTTTTAGT	1440
AAGACAAAGG AGAAAATAGA TGATTTATGC AGGAATTCTT GCCGGTGGAA CTGGCACACG	1500
CATGGGGATC AGTAACCTGC CAAAACAATT TTTAGAGCTA GGTGATCGAC CTATTTGAT	1560
TCATACAAATT GAAAATTIG TCTTGGAGCC AAGTATTGAA AAAATTGTAG TTGGTGTCA	1620
TGGAGACTGG GTTTCTCATG CAGAAGATCT TGTAGATAAA TATCTCCTC TTTATAAGGA	1680
ACGTATCATC ATTACAAAGG GTGGTGTGA CCGCAATACA AGTATTAAGA ACATCATTGA	1740
AGCCATTGAT GCTTATCGTC CGCTTACTCC AGAGGATATC GTTGTACCC ACGATTCTGT	1800
TCGTCCATT ATTACACTTC GCATGATTCA GGACAATATC CAACTTGCCTC AAAATCATGA	1860
CGCAGTGGAC ACAGTGGTAG AAGCGGTGA TACTATCGTT GAAAGTACCA ATGGTCAATT	1920
TATTACAGAT ATTCCAAATC GTGCTCACCT TTATCAAGGA CAAACACCTC AAACATTCCG	1980
TTGCAAGGAC TTCATGGACC TTTATGGATC TCTTCTGAT GAAGAGAAGG AAATCTTGAC	2040
AGATGCATGT AAAATCTTG TGATCAAAGG AAAAGATGTG GCTTGGCCA AAGGTGAATA	2100
CTCAAATCTG AAGATTACAA CCGTAACAGA TTTGAAGATT GCAAAAGTA TGATTGAGAA	2160
AGACTAGTAA AATGATTAAT CAAATTATC AACTAACTAA GCCTAAGTTT ATCAATGTCA	2220
AATATCAGGA AGAGGCTATT GACCAAGAGA ATCATATCCT TATCCGTCCC AACTACATGG	2280
CTGTCTGTCA TGCGGATCAG CGTTACTATC AGGGAAAACG TGATCCCAAG ATTTTGAAATA	2340
AAAAGCTTCC AATGGCAATG ATTACAGAGT CATGTGGAAC CGTCATTCT GACCCGACCG	2400

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GAACCTACGA GGTTGGTCAA AAAGTTGTCA TGATTCCCAA TCAGTCTCCT ATGCAGAGTG	2460
ATGAAGAAATT CTATGAAAAC TACATGACAG GGACCCATT CTTGTCTAGT GGATTGATG	2520
GCTTTATGAG AGAGTTGTT TCTCTCCCTA AAGATCGTGT GGTGGCTTAT GATGCTATTG	2580
AAGATAACGGT TGCGACCCATT ACAGAGTTG TCAGTGTGGG CATGCACGCT ATGAATCGTC	2640
TATTGACTCT TGCTCATAGC AAGCGGGAGC GGATCGCCGT TATTGGAGAT GGAAGTTAG	2700
CTTTGTGGT TGCCAATATT ATCAACTATA CTTTGCCAGA AGCAGAGATT GTGGTTATTG	2760
GTCGTCATTG GGAAAAGTTG GAACTCTTCT CATTGCCAA AGAATGCTAT ATTACGGATA	2820
ATATTCCCTGA AGATTGGCC TTTGACCATG CTTTTGAATG TTGTGGTGGT GATGGTACTG	2880
GACCAGCTAT TAATGACTTG ATTCGCTACA TTCGTCCTCA GGGAACGATT CTCATGATGG	2940
GAGTTAGCGA ATATAAAGTC AATCTCAATA CTCGCGATGC CTTAGAAAAG GGCTTGATTT	3000
TGGTTGGGTC ATCTCGTTCT GGTCGCATTG ATTTTGAAAA TGCTATCCAA ATGATGGAAG	3060
TCAAGAAATT TGCCAATCGT CTTAAAATA TCCTTTATCT AGAAGAACCT GTAAGAGAAA	3120
TTAAAGATAT TCATCGTGTCT TTTGCAACCG ATTTAACAC AGCCTTAAA ACAGTGTAA	3180
AGTGGGAAGT ATAAGTACTG GAGGTTAATT GTGGAGAAAA TCATTAAGA AAAAATTCT	3240
TCCTTACTTA GTCAAGAAGA GGAAGTCCTC AGTGTGAAC AACTGGGTGG AATGACCAAT	3300
CAAAACTATT TGGCCAAAAC AACAAATAAG CAATACATTG TTAAATTCTT TGGTAAAGGG	3360
ACAGAAAAGC TTATCAATCG ACAAGATGAA AAGTACAATC TTGAACTACT AAAGGATTAA	3420
GGCTTAGATG TAAAAAATTAA TCTTTTGAT ATTGAAGCTG GTATCAAAGT AAATGAGTAT	3480
ATCGAATCTG CGATTACGCT TGATTCAACG TCAATCAAGA CCAAGTTCGA CAAAATTACT	3540
CCAATATTAC AACTATTCA TACGCTCGCT AAGGAATTAA GAGGAGAATT TGCTCCTTT	3600
GAAGAAATCA AAAAATACGA ATCCTTGATT GAAGAACAAA TTCCTTATGC CAACTATGAA	3660
TCTGTTAGAA ATGCAGTCTT CTCCTTAGAG AAAAGACTGG CTGACTTAGG TGTTGACAGA	3720
AAATCTTGTC ATATCGATTT GGTGCCTGAA AACTTTATCG AATCACCTCA AGGACGACTT	3780
TATTTGATG ACTGGGAATA TTCATCAATG AATGATCCAA TGTGGGATTT GGCTGCCCTC	3840
TTTTTAGAGT CTGAATTCACT TTCCCAAGAG GAAGAAACTT TCTTATCTCA CTATGAGAGT	3900
GACCAAAACAC CGGTTCTCA TGAAAAGATT GCTATTATA AAATTTACA AGATACTATT	3960
TGGAGTCTAT GGACTGTCTA TAAGGAAGAG CAAGGTGAAG ATTTGGTGA CTATGGTGTG	4020
AATCGTTACC AAAGAGCTAT TAAAGGTTG GCTTCTTATG GAGGTTCAAGA TGAAAAGTAA	4080
AAACGGAGTT CCTTTGGCC TTCTCTCAGG TATTTCTGG GGCTTGGGTC TAACGGTTAG	4140
TGCTTATATC TTTTCGATTT TTACAGATTG GTCACCCATT GTGGTGGCTG CAACTCATGA	4200

817

TTTTTGAGC ATCTTATCT TACTAGCTT TCTCTGGTA AAAGAAGGGA AAGTCGCCT	4260
CTCAATTTC TTAAATATTC GCAATGTCAG TGTTATCATC GGAGCCTG TAGCAGGCC	4320
TATCGGTATG CAGGCCAATC TTTATGCAGT TAAGTATATC GGAAGTCTT TAGCTTCATC	4380
TGTATCGGCT ATTTACCCCTG CGATTCAGT TCTATTGGCT TTCTTCTTT TGAAGCACAA	4440
GATTCGAAA AATACTGTAT TTGGGATTGT CTTGATTATT GGAGGGATTA TTGCTCAGAC	4500
CTATAAGGTT GAACAGGTT ATTCTTTCTA CATTGGGATT CTTTGTGCTT TGGTTGTGC	4560
TATTGCATGG GGAAGTGAGA GTGTTCTTAG CTCTTTGCC ATGGAAAGTG AATTGAGTGA	4620
AATCGAAGCC CTCTTAATCC GTCAAGTAAC TTCGTTCTTG TCCTATCTTG TGATTGTGCT	4680
CTTCTCTCAT CAGTCATTAA CTGCAGTAGC CAATGGACAA TTGCTAGGTC TCATGATTGT	4740
TTTTGCAGCC TTTGATATGA TTTCCTACTT GGCTTATTAT ATCGCTATCA ATCGCTTGCA	4800
ACCAGCCAAG GCTACAGGCT TGAACGTGAG CTATGTAGTA TGGACGGTCT TGTGTTGCAGT	4860
TGTTTCTTG GGTGACCCGC TAGATATGCT GACCATTATG ACGTCACTTG TCGTCATTGC	4920
TGGAGTTTAT ATTATTATTA AAGAATAAAG GAGATTGCG TGAAAGCCAT TATCTTAGCA	4980
CGGGGATTGG GAACTCGCTT GCGTCCTATG ACTGAAAATA CCCCTAAAGC CTTGGTTCAG	5040
GTAAATCAA AAACTTTGAT TGAGTACCAA ATTGAGTTTC TCAAAGAAAA AGGAATCAAT	5100
GACATCATCA TCATTGTTGG TTATCTAAA GAACAATTG ATTACTGAA AGAGAAATAC	5160
GGTGTTCGTC TCGTTTCAA TGATAAACAC GCTGACTACA ATAACCTTTA CTCTCTCTAT	5220
CTTGTAAAAG AAGAATTGGC CAACAGCTAT GTTATTGATG CTGACAATTA TCTCTTTAAA	5280
AATATGTTCC GCAATGATTT GACACGTTCG ACTTATTTA GTGTTATCG TGAAGATTGT	5340
ACCAACGAAT GGTCTTGGT TTATGGAGAT GACTACAAGG TTCAAGACAT TATTGTTGAT	5400
AGCAAGGCAG GTCGCATCCT TAGTGGGTGA TCCTTCTGGG ATGCTCCAAC TGCAGAAAAG	5460
ATTGTCAGCT TTATCGACAA GGCTTATGTA AGTGGTGAAT TTGTTGATCT CTATTGGGAC	5520
AATATGGTTA AGGATAATAT CAAAGAGCTA GATGTCTATG TTGAAGAATT AGAAGGCAAT	5580
AGCATTATG AGATCGATAG TGTCCAAGAC TATCGTAAAT TAGAAGAAAT TCTTAAAAAC	5640
GAAAATTAAA GATTCCAACA TCTGACAAAA TAGTCGGATG TTTTTGATT TTTTACGAAC	5700
TTTTACGAAT AGATAGATGA GTAGAAAAAG AAATGGAGTT ATTTATGAAA ATCACAAACT	5760
ATGAAATCTA TAAGTTAAA AAATCAGGTT TGACCAATCA ACAGATTGTT AAAGTGCTAG	5820
AATAACGGTGA AAATGTTGAT CAGGAGCTTT TGTTGGGTGA TATTGCAGAT ATCTCAGGTT	5880
GCGTAATCC AGCCGTTTT ATGGAACGTT ATTTTCAGAT AGACGATGCG CATTGTCGA	5940

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AAGAGTTCA AAAATTCCA TCTTCCTCTA TTTAGATGA CTGTTATCCT	TGGGATTGAA	6000
GTGAAATATA TGATGCCCT GTACTTTAT TTTACAAGGG AAATCTTGAC	CTCCTGAAAT	6060
TCCCGAAGGT AGCGGTCGTG GGCAGTCGTG CTTGTAGCAA ACAGGGAGCT	AAGTCAGTTG	6120
AAAAAGTCAT TCAAGGCTTG GAAAATGAAC TGGTTATTGT CAGTGGTCTG	GCCAAGGGCA	6180
TTGACACAGC AGCTCATATG GCAGCTCTTC AGAATGGCGG AAAAACCAATT	GCAGTGATTG	6240
GAACAGGACT GGATGTGTTT TATCCTAAAG CCAATAAACG CTTGCAAGAC	TACATCGGCA	6300
ATGACCATCT GGTCTAAAGT GAATATGGAC CTGGTGAACA ACCTCTGAAA	TTTCATTTTC	6360
CTGCCGTAA TCGCATCATT GCTGGACTTT GTCGTGGTGT GATTGTAGCA	GAGGCTAAGA	6420
TGGGTTCAAG TAGTCTCATT ACGTGTGAGC GAGCAATGGA AGAAGGACGC	GATGTCCTTG	6480
CTATTCCCTGG TAGCATTAA GATGGACTAT CAGACGGTTG CCATCATTG	ATTCAAGAAG	6540
GAGCAAAATT GGTCACCAGT GGGCAAGATG TTCTTGCGGA ATTTGAATTT	AAAAATGAC	6600
CTAAGCTAGA ATTCTAAGAA AAAATCAATT TTAAGAGAAA ATGAACCCAA	CATTTCCATA	6660
ATAAAACGCA TATTAGCAAG TTTTAACAC TTGATAATAT GCGTTTTTC	TAAGTGGATT	6720
AGTAGAGTAG AGGATTTTC TCATATAATA CTCTTCGAAA ATCTCTCAA	ACTACGTCAG	6780
CTTCCATCTG CAACCTCAAA ACAGTATTTT GAGCgATTC GTCAGTCTTA	TCTACAACCT	6840
CAAAGCAGTG CTTTGAGCAA CCTGTGGCTA GCTTCCTAGT TTGCGCTTTG	ATTTTCATTG	6900
AGTATAAGGG AAAGTATAGT GAATTGAAAT AAGATGTGAA CAACTCTATC	AGGAAAGTCA	6960
AATTAATTAA TAGAAATATT TTAGCAGCCA AGGTGTACTG TTATAGATTC	AATTACACTA	7020
TAATTTAGTG TAATTGAGAA AGGAGAAATG ATTGTGATTG ATGTTGGCTA	GGTTATGTT	7080
AATGATTCTT ACCGTCTCAA ATCTTGTCAAG TAAGGAAAAA TAAATTCTTC	AAAAGTAGAG	7140
ATTACAAGGC TTGTTTAAGA AAGAATTCAA AGACCTTGAC AAATAAAAT	AAAATGGTTA	7200
TTATAAAAAA TGGTCTGAAA TAGATGATGA TACTTTCGA AAATCTCTTC	AAATACGTCA	7260
GCTCAGCTTT GCCTTGCTGT GTTTGAGCA AGCTACGGTT AGCTTCCGAG	TTTGATTT	7320
ATTTACTAGA AATGAAACTG ATGAGAGATA TCAGTAGACAA TTGAGTCAG	GATATTATGG	7380
AAAATGATAA AAAGAGCTCG TGAGATTGGC ATATCAGACT ACTAAAGTAT	TGAGTTGTT	7440
AGGATTTAG CGACTAGTTA GCTGGGAAAG GAAGATATTG GTGACAAATA	ATAAACTGTA	7500
TTCGTTGATA GAATTAGAA ATAAAATATA TGAAGAATTAA GAACTTCCA	GAAGTGATT	7560
AGCGATTTA CTATGTGCCA TGCTTATCGC CTCTATCGGA TTAAATATGG	ATTCGACTCC	7620
CGTGATTATT GGAGCCATGT TAATCTCTCC TTTGATGACA CCTATTCTGG	GAGTGGGGCT	7680
CTCTCTAGCT ATATTTGATT TTAAATTGTT AAGAAAATCT TTTAAAATAT	TAGCTATTCA	7740

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AATTCTTGCC	AGTCTAATAG	CTTCAACACT	TTATTTTAT	CTTTCTCCA	TTTCGTATGC	7800
TAGTTCGGAG	ATTGTTGCTA	GAACCTCTCC	GACTATTTGG	GATGTTCTCA	TTGCTTTGT	7860
AGGAGGGATA	GCAGGTATCA	TTGGTGCTAG	GAAAAAAAGAG	AC		7902

(2) INFORMATION FOR SEQ ID NO: 113:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18627 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 113:

GAAGTTGAAA	TGGCCAGCTG	ATGAGCAATA	TCGGTCATAG	AAATCTTCTC	AATCAACTTT	60
TGCGCAATT	TTTGGTGAT	AATAACGAGGA	ATTTGGTGAT	TTTTCTTGAC	GATAGAAGTT	120
TCAGCGACCA	TCATTTTGTA	ACAGTGATAG	CACTTGAAAC	GACGCTTCT	AAGTAGAATT	180
CTAGTAGGCA	TACCAGTTGT	CTCAAGGTAA	GGAATCTTAG	ACGGTTTTG	AAAGTCATAT	240
TTCTTCATT	GGTTTCCGCA	CTCAGGGCAA	GATGGGGCGT	CGTAGTCCAG	TTTGGCGATG	300
ATTTCCCTGT	GTGTATCTT	ATTGATGATG	TCTAAAATCT	GGATATTAGG	GTCTTTAATG	360
TCTAGTAATT	TTGTGATAAA	ATGTAATTGT	TCCATATGAA	TCTTTCTAAT	GAGTTGTTG	420
GTCGCTTTC	ATTATAGGTC	ATATGGGACT	TTTTTCTAC	AATAAAATAG	GCTCCATAAT	480
ATCTATAAGG	GATTTACCCA	CTACAAATAT	TATAGAGCCA	AAAATCCTT	GTTTACTAAA	540
CAAGGGATTT	TTCTTTGTC	TCTGCTCCTT	TTTGATATA	ATAGTTCTAT	GTTAAAATCA	600
GAAAAACAAT	CACGTTATCA	AATGTTAAAT	GAAGAATTGT	CCTTCCTATT	GGAAGGCGAA	660
ACCAATGTT	TGGCTAATCT	TTCCAACGCC	AGTGCTCTCA	AAAATCACG	TTTCCTTAAT	720
ACCGTATTG	CAGGCTTTA	TTTGTTCGAT	GGAAAGGAAT	TGGTTTAGG	CCCCTCCAA	780
GGAGGTGTT	CCTGCATCCG	TATTGCACTA	GGCAAGGGTG	TTTGTGGTGA	GGCAGCTCAC	840
TTTCAGGAAA	CTGTTATTGT	TGGAGATGTG	ACGACCTATC	TCAACTATAT	TTCTTGTGAT	900
AGTCTAGCTA	AAAGTGAAT	TGTGGTCCG	ATGATGAAGA	ATGGTCAGTT	ACTTGGAGTT	960
CTGGATCTGG	ATTCTTCAGA	GATTGAGGAT	TACGATGCTA	TGGATCGAGA	TTATTTGGAA	1020
CAATTGTCG	CTATTTGCT	TGAAAAGACA	GCATGGGACT	TTACGATGTT	TGAGGAAAAA	1080
TCTTAATGTA	TCAAGCACTT	TATCGAAAAT	ATAGAAGTCA	AAACTCTCC	CAGTTAGTTG	1140
GTCAAGAAGT	TGTGGCTAAG	ACTCTAAC	AAGCGGTGGA	GCAAGAGAAA	ATAAGTCACG	1200

820	
CTTATCTTT TTCTGGTCCT CGTGGAACGG GAAAAACCAG TGTTGCTAAA ATCTTGC	1260
AGGCTATGAA CTGTCCCAAT CAAGTGGGTG GCGAACCTTG CAATAACTGC TATATTGTC	1320
AAGCAGTGAC GGACGGTAGT TTAGAAGATG TCATTGAAAT GGATGCAGCT TCTAATAATG	1380
GGGTAGATGA AATTCGCAA ATTCTGTATA AATCTACCTA TGCGCCTAGC CTTGCTCGTT	1440
ATAAGGTTTA TATCATAGAT GAGGTTCACAA TGCTGTCTAC AGGGGCTTT AATGCCCTCC	1500
TAAAGACGCT GGAAGAACCA ACACAGAATG TAGCTTTAT TTTGGCCACT ACTGAATTGC	1560
ACAAGATTCC TGCTACTATT CTATCCCGTG TGCAACGTT TGAGTTAAA TCAATTAAAGA	1620
CACAGGATAT TAAGGAACAT ATTCACTATA TCTTAGAAAA AGAAAATATC AGTTCTGAAC	1680
CAGAGGCTGT GGAAATCATT GCCAGACGGG CGGAAGGTGG AATGCGGGAC GCCTTGTCTA	1740
TTTGGATCA AGCCCTGAGT TTGACACAGG GAAATGAGCT GACGACTGCT ATCTCTGAAG	1800
AAATTACTGG CACCATTAGC CTATCAGCCT TGGATGATTA TGTGGCGGCC TTGTCCTAAC	1860
AGGATGTTCC CAAAGCTTG TCTTGCTTGA ATCTTCTTT TGACAATGGT AAGAGCATGA	1920
CTCGTTTGTG GACCGATCTT TTGCACTATT TAAGAGACTT GTTAATTGTT CAAACAGGGG	1980
GAGCAAATAC TCATCATAGT TCAGTCTTG TAGAAAATTG GGCACCTCCT CAAAAAAATC	2040
TGTTTGAAAT GATTGCTTA GCAACAGTGA GTT TAGCAGA TATTAAGTCT AGTTGCAAC	2100
CCAAGATTAA TGCTGAAATG ATGACCGTCC GTTGGCGGA AATCAAGTCC GAACCAGCTC	2160
TATCAGGAGC GGTTGAAAAT GAAATTGCTA CGCTGAGACA GGAAGTTGCC CGTCTCAAAC	2220
AAGAGCTTTC TAATGTAGGT GCGGTTCTA AACAAAGTTGC ACCAGCTCCT AGTCGACCAG	2280
CTACGGGCAA AACAGTCTAT CGTGTGATC GCAATAAGT GCAATCTATC TTACAAGAGG	2340
CCGTCGAAAA TCCTGATTTA GCACGTCAAATTTAATTCG TTTGCAGAAT GCCTGGGGAG	2400
AGGTAATTGA AAGTCTAGGT GGGCCGGACA AGGCTCTGCT AGTTGGTTCT CAACCGGTTG	2460
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TGAAACGAGA CAATCTCAAT ACCATGTTG GTAATATCCT CAGTCAGGCG GCAGGTTTT	2580
CACCTGAGAT TTTAGCTATT TCCATGGAGG AATGGAAAGA AGTTGCGCAGA GCCTTTTCAG	2640
CCAAAGCCAA ATCTTCTCAA ACTGAAAAG AAGTAGAAGA AAGCCTGATT CCAGAAGGAT	2700
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TACTCTTTAG AAATTCGGA GTCAGTTATG TGATGGCGAA AATCGTTGAT GTCATCGATC	2940
AGCATTAA TAGGAAAGAC TAGCCCTCAG CTTCCAGACA AAATCAAAGC CTTTTAGGCT	3000

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TTTTTTGTT ATACTAGAAA AGTATATTAA TAGAATTTC GCTCTATTTC TGGGGAAATC	3060
AGACGTTTT CTAGTAAGTA CTGTAAAAGT TTTGAAAAAG AAAGGAACTA TCATGTCAGT	3120
ATTAGAGATC AAAGATCTTC ACGBTGAGAT TGAAGGAAAA GAAATTTAA AAGGGTTAA	3180
CCTGACCCTG AAAACAGGAG AAATTGCCGC TATCATGGGA CCAAATGGTA CAGGTAATC	3240
GACTCTTCT GCCGCTATCA TGGGAAATCC AAACATGAA GTAACTAAAG GTGAAGTTT	3300
GTGGATGGC GTAAACATCC TTGAGTTGGA AGTGGATGAG CGTGCCTGTA TGGGACTTT	3360
CCTTGCTATG CAATACCCAT CAGAAATCCC TGGATTACCA ATGCTGAGT TTCTCGTGC	3420
CGCTATGAAT GCGGGTAAAG AAGATGATGA GAAGATTCA GTTCGTGAGT TTATTACTAA	3480
GCTAGATGAA AAAATGGAAT TGCTAACAT GAAAGAAGAA ATGGCAGAGC GTTACCTCAA	3540
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TGTGTCTAA GGTGTCAATG CCATGCGTGG TGAAGGTTTT GGTGCTATGA TCATCACTCA	3720
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TGTTGTCCTT TCTGGTGGTC CAGAATTGGC TGCCTGTTG GAACGTGAAG GATAACGAAA	3840
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GCCATCAGCA AATGTTCCAG ATTTCACAGC TTTAGATCAT CACTTGAAAGT TGGTCAAGT	4140
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TGCAAGTATT GACTGGGCTA TCGGTGTCAT GAACGAAGGA AATGTCGTTG CTGATTTGA	4680
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CATCATCAAG	GGTGCTAAGG	GAGCAGATGC	GCAACAAGAG	AGCCGTGTTC	TCATGCTTTC	4920
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CTTTATATGG	CAGTGGTAGC	AGACCATTG	AAAAATCCAC	ATCACCAAGG	GAAGTTAGAA	6480
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CCTTTACAAA AGAACTTCCA ATGGAATACG CAGTTGAGCT GAACCGCTTG ATTAGCTATG	8280

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ACGTCAGCAT CGCCTTACCG TATGTATGGT TwCTGATTCG TCAGTTTCAT CTACAACCTC	8400
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GTTGGCAGGT	GTTTGGACGT	GGGTTCGACT	CCCACCGGCT	CCATTATTCC	TTTGCATTCT	11880
TTTGCATTCC	TTGGTAAAC	GTTGTTAAAT	CAACGTTTT	TATTTTATC	TTTGGTATTTC	11940
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GGAGCAATCG	TAAGTGGTGT	GGTTTATGGT	TACCTACGCA	AACCACAAGC	ATAAAAAAATA	15900
GAAAAATGAA	AAGATTGGAC	CGTTTGGTGC	AGTCTTTTC	TCTTCCCAGA	ATGCCTGTGA	15960
AATATGGTAT	AATAGAAGAA	TGGCAAACAA	GAATACAAGT	ACAACAAGAC	GGAGACCGTC	16020
TAAAGCAGAA	CTGGAAAGAA	AAGAACGAT	TCAACGAATG	TTGATTTCGT	TAGGAATTGC	16080
GATTTTATTG	ATTTTCCGAG	CCTTCAAATT	AGGGGCTGCA	GGTATAACCC	TTTATAATTT	16140
AATTCGCTTG	CTAGTGGTA	GCCTAGCTTA	TCTGGCGATA	TTCGGCTAT	TAATCTATCT	16200
CTTCTTTTC	AAAGTGGATAC	GAAAACAGGA	AGGACTCTTA	TCTGGCTTT	TCACCATATT	16260
TGCTGGCTTA	CTCTTGATTT	TTGAGGCCTA	CTTGGTTTGG	AAATATGGTT	TGGACAAGTC	16320
CGTTCTAAA	GGGACCATGG	CTCAGGTTGT	GACAGATCTG	ACTGGTTTC	GAACGACTAG	16380
CTTTGCTGGA	GGGGGCTTGA	TCGGGGTCGC	TCTTTATATT	CCAACAGCCT	TTCTCTTTTC	16440
AAATATCGGA	ACTTACTTTA	TTGGTTCTAT	CTTGATTTA	GTGGGTTCTC	TCCTAGTCAG	16500
CCCTTGGTCT	GTTTACGATA	TTGCTGAATT	TTTCAGTAGA	GGCTTTGCCA	AATGGTGGGA	16560
AGGGCACGAG	CGTCGAAAAG	AGGAACGCTT	TGTCAAACAA	GAAGAAAAAG	CTCGCCAAA	16620
GGCTGAGAAA	GAGGCTAGAT	TAGAACAAAGA	AGAGACTGAA	AAAGCCTTAC	TCGATTTGCC	16680
TCCTGTTGAT	ATGGAAACGG	GTGAAATTCT	GACAGAGGAA	GCTGTTCAA	ATCTTCCACC	16740
TATTCCAGAA	GAAAAGTGGG	TGGAACCAGA	AATCATCCTG	CCTCAAGCTG	AACTAAATT	16800
CCCTGAACAG	GAAGATGACT	CAGATGACGA	AGATGTTCA	GTCGATTTT	CAGCCAAAGA	16860
AGCCCTTGAA	TACAAACTTC	CAAGCTTACA	ACTCTTGCA	CCAGATAAAC	CAAAGATCA	16920
GTCTAAAGAG	AAGAAAATTG	TCAGAGAAAA	TATCAAAATC	TTAGAAGCAA	CCTTGCTAG	16980
CTTTGGTATT	AAGGTAACAG	TTGAACGGGC	CGAAATTGGG	CCATCAGTGA	CCAAGTATGA	17040
AGTCAAGCCG	GCTGTTGGTG	TAAGGGTCAA	CCGCATTCC	AATCTATCAG	ATGACCTCGC	17100
TCTAGCCTTG	GCTGCCAAAG	ATGTCCGGAT	TGAAGCACCA	ATCCCTGGGA	AATCCCTAAT	17160

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CGGAATTGAA	GTGCCCAACT	CCGATATTGC	CACTGTATCT	TTCCGAGAAC	TATGGGAACA	17220
ATCGCAAACG	AAAGCAGAAA	ATTCCTTGGA	AATTCCCTTA	GGGAAGGCTG	TTAATGGAAC	17280
CGCAAGAGCT	TTTGACCTTT	CTAAAATGCC	CCACTGCTA	GTTGCAGGTT	CAACGGGTTC	17340
AGGGAAGTCA	GTAGCAGTTA	ACGGCATTAT	TGCTAGCATT	CTCATGAAGG	CGAGACCAGA	17400
TCAAGTTAAA	TTTATGATGG	TCGATCCCAA	GATGGTTGAG	TTATCTGTTT	ACAATGATAT	17460
TCCCCACCTC	TTGATTCCAG	TCGTGACCAA	TCCACGCAA	GCCAGCAAGG	CTCTGCAAAA	17520
GGTTGTGGAT	GAAATGGAAA	ACCGTTATGA	ACTCTTGCC	AAGGTGGGAG	TTCGGAATAT	17580
TGCAGGTTTT	AATGCCAAGG	TAGAAGAGTT	CAATTCCCAG	TCTGAGTACA	AGCAAATTCC	17640
GCTACCATTC	ATTGTCGTGA	TTGTGGATGA	GTTGGCTGAC	CTCATGATGG	TGGCCAGCAA	17700
GGAAAGTGGAA	GATGCTATCA	TCCGCTTGG	GCAGAAGGCG	CGTGCTGCAG	GTATCCACAT	17760
GATTCTTGCA	ACTCAGCGTC	CATCTGTTGA	TGTCATCTCT	GGTTTGATTA	AGGCCAATGT	17820
TCCATCTCGT	GTAGCATTG	CGGTTTCATC	AGGAACAGAC	TCCCCTACGA	TTTTGGATGA	17880
AAATGGAGCA	GAAAAACTTC	TTGGTCGAGG	AGACATGCTC	TTTAAACCGA	TTGATGAAAA	17940
TCATCCAGTT	CGTCTCCAAG	GCTCCTTAT	CTCGGATGAC	GATGTTGAGC	GCATTGTGAA	18000
CTTCATCAAG	ACTCAGGCAG	ATGCAGACTA	CGATGAGAGT	TTTGATCCAG	GTGAGGTTTC	18060
TGAAAATGAA	GGAGAATTTC	CGGATGGAGA	TGCTGGTGGT	GATCCGCTTT	TTGAAGAAGC	18120
TAAGTCTTTG	GTTATCGAAA	CACAGAAAGC	CAGTGCCTCT	ATGATTTCAGC	GTCGTTTATC	18180
AGTTGGATTT	AACC GTGCAGA	CCC GTCTCAT	GGAAGAACTG	GAGATAGCAG	GTGTCATCGG	18240
TCCAGCTGAA	GGTACCAAAC	CTCGAAAAGT	GTTACAACAA	TAAAAAAATA	GCTTCTTCC	18300
AAGTTGGAG	GGAAAGCTATT	TTAGTGGCTA	TTGATTGCTT	TTATTTCTG	AAGTTGGCGC	18360
ATTGGACTGT	TTTTCGTTTT	CAGTAGCAGG	TTTACTTGAA	GCAGGAGTAG	AAGAGTCCCTG	18420
AGTTGCTGTT	TTCTGATCTT	CTTTTTCTC	TTCCCTGACG	CTAGATTTTG	GTGTTCCCTC	18480
TTGCTGTGTT	TTTCCTTGAC	TAGTGTAGT	CTCTTTAGTT	GGACTGGTGT	TTTCCTTAGG	18540
GGATTCCCTT	TGGATTTCTT	TGACAATGGT	TGTCGTCTGG	CTTGTGCTAG	GTTCTTTTT	18600
AATATTTTG	TTATTATCCA	AGGCGTT				18627

(2) INFORMATION FOR SEQ ID NO: 114:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 2560 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 114:

TAAAATACGT	TACCTTGCTT	CTGCACGTT	AGCAGGTAAG	TCATTGAAAT	TTAAAGATCA	60
AGATATTACA	ATTGAAGAAA	CGACTGAAAC	AGCTTTGAA	GGAGTTGATA	TTGCTCTCTT	120
TTCAGCAGGT	AGTTCTACAT	CAGCTAAGTA	TGCACCATA	GCAGTAAAAG	CTGGCGTGGT	180
AGTAGTAGAT	AATACATCTT	ATTTCCGTCA	AAATCCAGAT	GTTCCTTGG	TTGTTCCAGA	240
GGTCAATGCT	CATGCACTTG	ATGCTCACAA	CGGAATCATT	GCCTGCCCTA	ATTGTTCAAC	300
AATTCAAATG	ATGGTGGCTC	TTGAGCCGGT	TCGCCAAAAA	TGGGGCTTGG	ACCGTATCAT	360
TGTTTCAACT	TATCAAGCCG	TTTCAGGTGC	TGGTATGGGA	GCAATTCTTG	AGACACAACG	420
TGAACCTCGT	GAAGTCTTGA	ATGATGGTGT	GAAACCACGT	GATTTGCATG	CGGAAATCTT	480
GCCTTCAGGT	GGTGACAAGA	AACATTATCC	TATCGCCTTT	AACGCTTTC	CACAAATTGA	540
TGTTTCACT	GATAATGATT	ACACGTACGA	AGAGATGAAG	ATGACCAAGG	AAACTAAGAA	600
AATTATGGAA	GATGATAGCA	TTGCACTATC	TGCAACATGT	GTGCGTATTC	CAGTCTTGT	660
AGCTCACTCT	GAGTCTGTTT	ATATCGAAAC	AAAAGAAGTG	GCTCCAATCG	AAGAAGTAAA	720
AGCAGCTATC	GCAGCCTTCC	CAGGTGCTGT	TCTTGAAGAT	GATGTAGCTC	ATCAAATCTA	780
TCCTCAAGCT	ATCAATGCAG	TTGGTTCGCG	TGATACCTTT	GTTGGTCGTA	TCCGTAAAGA	840
CTTGGATGCA	AAAAAAGGAA	TTCACATGTG	GGTTGTTCA	GATAACCTTC	TCAAAGGTGC	900
TGCTTGGAAC	TCAGTTCAGA	TTGCTGAAAC	TCTTCATGAA	CGTGGATTGG	TTCGTCCAAC	960
AGCCGAATTG	AAATTTGAAT	TAAAATAGTC	ATATCGTTA	GGAGTTCAGA	TGAACTCCTT	1020
CTTTGAAATA	GAGAGGTGTT	TTCGTGTCTT	ATCAAGATT	AAAAAAATGT	AAAATCATTA	1080
CAGCCTTTAT	TACCCCCCTTC	CATGAGGATG	GTTCCATTAA	CTTGATGCT	ATTCCAGCCT	1140
TGATTGAGCA	TTTATTGGCC	CATCATACGG	ATGGAATTCT	TCTCGCAGGA	ACGACTGCTG	1200
AGAGTCCAAC	TTTGACCCAC	GATGAGGAGT	TGGAGTTGTT	TGCGGCTGTA	CAAAGGTTG	1260
TCAATGGACG	CGTTCCCTTG	ATTGGGGGTG	TAGGTACTAA	TGATACGCGT	GACTCTATTG	1320
AGTTTGTCAA	AGAAGTAGCG	GAATTGGTG	GTTCGCGAGC	TGGGCTTGCT	ATTGTTCCCTT	1380
ACTACAACAA	ACCTTCTCAA	GAAGGGATGT	ATCAGCACTT	TAAGACTATT	GCAGATGCTT	1440
CTGACCTACC	AATTATTATC	TATAACATTC	CAGGGCGTGT	AGTTGTCGAA	TTGACTCCAG	1500
AAACCATGCT	TCGCTTGGCT	GACCATCCAA	ATATTATCGG	TGTCAAAGAA	TGTACTAGCT	1560
TGGCTAATAT	GGCTTACTTG	ATTGAGCACA	AGCCTGAAGA	GTTCTTGATT	TATACAGGTG	1620
AGGATGGAGA	TGCTTTCCAT	GCCATGAACC	TTGGGGCGGA	TGGGGTTATT	TCTGTTGCCT	1680

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CTCATACAAA	TGGGGATGAA	ATGCACGAGA	TGTTTACTGC	GATTGCAGAA	AGCGATATGA	1740
AGAAAGCCGC	AGCAATTCA	CGTAAATTCA	TTCCTAACGGT	TAATGCTCTC	TTCTCTTATC	1800
CAAGTCTGC	TCCAGTTAACG	GCAATTCTTA	ACTATATGGG	ATTTGAAGCT	GGACCCACTC	1860
GTCTACCTCT	TGTTCCAGCA	CCAGAAGAAG	ATGCCAACG	CATTATCAAG	GTTGTCGTAG	1920
ATGGCGACTA	CGAAGCAACT	AAGGCAACTG	TAACAGGGT	CTTAAGACCA	GATTACTAAT	1980
AAAGACAATA	AAATCCGGCT	CTTTGTCAAC	TGTAGTGGGT	TGAAGTCAGC	TAAGCTCGAG	2040
AAAGGACAAA	TTTTGTCTT	TCTTTTTGTA	TATTCAGAGC	GATAAAAATC	CGTTTTTGTA	2100
AGTTTTCAA	GTTCCGAAA	CCAAAGGCAT	TGCGCTTGAT	AAGTTTGATG	AGATTATTGG	2160
TCGCTTCCAA	TTTGGCGTTT	GAATAGGGTA	GTTGAAGGGT	GTTGACGATT	TTCTTTTGTT	2220
CCTTTAGAAA	GGTTTTAAAG	ACAGTCTGAA	AAATAGGATG	AACCTGCTTC	AGATTGTCCT	2280
CAATGAGTCC	GAAAAATTTC	TCCGGTTCCCT	TATTCTGAAA	GTGAAACACC	AAGAGTTGAT	2340
AGAGCTGATA	GTGATGTTTC	AAGTTTGATG	AATAGCTCAA	AAGCTTGTTT	AAAATCTCTT	2400
TATTGGTTAA	GTGCATACGA	AAAGTAGGAC	GATAAAAATCG	CTTATCACTC	AGTTTACGGC	2460
TATCCTGTTG	AATGAGTTTC	CAGTAGCGCT	TGATAGCCTT	GTATTGGGA	TTTCGATGA	2520
AACTGATTCA	TGATTGGAC	ACGCACACGA	CTCATAGCAC			2560

(2) INFORMATION FOR SEQ ID NO: 115:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11303 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 115:

TATTGGATT	CCCTTGCAAT	CAGTTTATGG	GACAAGCACC	CGGCAGCGCA	GAGGAAATCA	60
ACGCCTTCTG	TAGCCTACAT	TTTCAAACCA	CCTTCCCACG	TTTGCCAAAG	ATTAAGGTCA	120
ACGGTAAGGA	AGCAGACCCCT	CTCTATGTCT	GGTTACAAGA	CCAGAAAATCC	GGCCCACTAG	180
GAAAACGAGT	CGAATGGAAT	TTCGCTAAGT	TTCTCATCGG	TCGAGATGGG	CAAGTCTTG	240
AACGCTTTTC	TTCAAAACAC	GACCCAAAAC	AAATTGAAGA	GGCGATACAA	ACTCTACTAT	300
AATTCAACAT	CTCACTATGA	TTAGGTTCC	TTAACCTGA	TGAATAGTGA	GATTTTTGTA	360
TGGGCTTTGA	CTTAAATAGA	AAAACACCCC	ATGATATGAA	ACATGAAGTG	TTGTAAAGTC	420
TATGTTGTAG	GTGCTTATT	CACAATTCA	ATGTGACCAG	TGATAACGAA	TACCATACAG	480

832	
AATCTTCATA TACACTAAC AAATGACTTT CTAATTATTT CAATTAGTTT TGGCTAGTAA	540
ATATCATTTC CAACAAACGC CCTCTCAATT CCTTATCCTG ATGATGCAAG ATATTCATTA	600
AGTCATGAGA GTTTTCGCA TTGATGAATT GATTTAACAA TCTATCTTT AATTCAATG	660
GAAGAGAACG TGTCTTAGT AGTCTAAAAA CTTCGTCATT TAAAGATGTC CTTTTATTAT	720
CTTTCCATTC AAATTTAGCT GTATCATTCT TATTTGGCAA TTCAATTATA GACACATTG	780
TTCCTTTAAA ATGAATTCTA TGTTTCTAT TGCTTGGAAC GATACTAGAA TCTCCTTGTA	840
ATGCTAACTC TACCATTCCC ATTTCCAAT CGATTGATAA TCTTGTTTA TATCTTGAC	900
CATTTTGATC TTCAAGCATT TCAAAAGAAT GTTGTTCAC TGGGAATACA TACCAATCTA	960
CAACTTCAGG TAAATCAACA CCCATACCTA TCTCAGAAC AACCAAGGGA ATGATTGCAC	1020
CACTTTTTGC AAACACAGGC GTAGTCGAGA TGTCCCTATA AACACTTAAC TTCACACCAC	1080
CTGTTGATTT TTTCTCTGAA AAGAAGTCAT ACCATTCACTTACAGGGAAC CATACTCTA	1140
CTTTTGAGA TTGGAATGTC AAATCCATCT TTTCTACAAT GGGAGGCCACC ATCAGTTCTG	1200
TTCCAAAAAA GTATTGGTTT GGAACATTAT AGCTCTCATC ATTCTCTGGA TAGAAATAAT	1260
AGATTGGACT GATTAATGGG GCACCTTCCT CATGTGTCTG TACATTCTG GTATATAGAT	1320
AGGGAATCAT CTGATGTCTC AAACGAAGGT ATTCTCTCAT AATCTTAGAT GTTGTTCCTG	1380
AAAAAAACCA AGGTTCTTTA CTATTAAGGACTCTAGA ACTATGTAAT CGAGTAATCG	1440
GACTAAAAAC ACCAAACTGT AGCCATCTAG TTTGTAGCTC TTGTCATAA TCCCCAACAA	1500
TATGTCCACC GATATCATGA CTCCACCAAC TATAACCGAT ATTAGATGCT GTCGCTGTAA	1560
AATAGGGTTG AAATCTTAAG GAATTCAAC TAATAATAGT ATCCCCTGAA AAACCAACAG	1620
GGTAGCGGTG ACTACCAGGA CCTGCATATC TTGATAAAAT CAAACCACCT TCTGCATTTT	1680
TACAACATATC CTGATAGTGA TAATGGTTA AAAGCCAAAG TGGATCTAGC ATACCTGTG	1740
TCCCTTGTG CCAGTCATC CACCAAAAT CTACTCCCTG CTTTTCTAGT TCATAATGAA	1800
CATCTTAAA GTAGGCTTCC CTAAAGAGG GATTAACAAA ATCAAAATA GCAGGTTCTT	1860
CTAGTTCTAC ATTTAACCC ACCGTTTG CGATTTGAGG ATAAGCTTCT TCATAAGCCC	1920
GTATCCCATC AGCAGGATGG ACATTTAAGG AGAGTTTAG CTTCTATCA TGAAAGTTGTT	1980
GCAATAACTG TTCTGGATTG GGTATTAAGT TTCTATTCCA ACTATATCCT GTCCAGGCCAC	2040
TTCCAAAGCG AGCTGGAATG TCAGTTATAT GCCAATCCAT ATCTAACACA CCGATAGATA	2100
ATGGAATTTCCTCTA AATCTGTCTA TTAAATCCAA GTATTCATCC GACGTATAAG	2160
GCCAATATCT ACTCCACCAA TTGCCTAAAG CATACTTTGG CAACAAGGGT GTTGAACCAG	2220
TCAAATGGTA AAAATCTCTG ATTGCTCCTC TATAATCATG CCCATAGGCA AAGAAATACA	2280

833

GGTCAATTTG ATTTCTCTC TCAATATAAC CAGATTGTC ATCCCAAATA AATCCTTGAG	2340
AATCATCCAA TAAGGCTATA CCATTTCGGC TAATAATTCC ATCTTCTAAC GAGATTGCTC	2400
CATCTGCCTT ATCCAGAGTC CGAGCTGTT CTTTTAACGT TTCAATAGAT TCACCAAAAT	2460
ACCAGCGACT ACCATATACG GCAAAATTTC CTTTTAATTC TATAAATAAA TTTTCGGCGT	2520
TAAAATTCTCC TTTATTAAAG TGCAGATGAA AATAGTCCGT CATAATATCT AGTACGTTG	2580
ATGTCCTCGAT ATAATCTAAC GAAATTTGGC CAAAATCTCT ATTATAGATA AGTTGTGTCG	2640
TTCTATCCTC AAAACTCCA GTTTGAGAGT ATTCTAACCT TACTAGCTT GCTGTTAATA	2700
CAGAGATTG ATAAAACTCT CCCTAAAAA TTTTCAATTG GTTTCCCTC TTTTATGGTA	2760
GCATAAAAAG AGAACGCACC ATTTTGATG CGTTTTCAT TATTCTGAAT GCAATGTTCT	2820
ATCTGTTATA TCTATGACAA ATAATAGTCA ATTGAAAAAA TGCACTGGAC AAAATATCTT	2880
TTAACAAACC AAGAGTTTAA TAAAGAGTTA TCACCTTCA ACTTTTCTAA GCTTATGCAG	2940
TTGTAACAAACA AACTACTTTT AAACATTAA CTAAGATAGG ATTGATAAAAT AATTCAAAAC	3000
TCTTACTAGC AATCATAACGA TATTCAAGCT CACGTGCTT TTTCCCTCCT GCTTATTCT	3060
TAGAACTGAA GAACCCGGAT CGGTATATAA ATTATCCGGA TCAACATAGT CATAAGATTC	3120
ATAACAGTTG CGCTTCATTA AGTCATCCCC AGAGCAAGAG CTTCATCTCG TAATTTTCA	3180
ACATCACTAA CCGTAGGTCG CCATCCTTCA ATCATATTG TACTTAAAGC ATACCAAACA	3240
CTCTTAAAAA CGGATCGGTT TTCAAAAGCT ATTCCCATGA TTGTCATCTT TTCTTATCT	3300
ATATCTAAGG ACATATGCTA CCTCCTTAG ATACATTATA CCATGTTCT CTGTAGCTT	3360
TAAAAATTAA ATTGTTGTTG TCATATCTAA GTTTTCAGCA CGCTTATCCT ATTTTATAAG	3420
CCTCAAACCC AAATATAAA CGCATTCTT TTGCTTTTT ACTATTGTAT CGTATTCTAC	3480
GATAACATAC TTACTTTAT TGTTTTTTA AATAACAGCA GTTCCCTGTT TATCAACTAT	3540
TCGAACTACT TTCTATTTG CTTCATACCC TACATAGCGA AAAATATGA AAAAGCAGAG	3600
AAGAATATCT TAAAAAGACC TCTTCACTGC TAATATTAAC ACTCATTATT TAAACTATAT	3660
GGATTCTATC ATCGAGTATA CTTTTTACT TATTAGATAC CTTGCTCTTC TTTCACCAAT	3720
TTTTGATCAT ATACACGGAT GAATGGAAGA TAGACTAGGA ATGCTGAAA TGACACATACT	3780
AGAGCAACTA ATACAGCTCG AAGATCTGCT GTCCCTAAGA AAGCTCCAAT CCCTACTGGA	3840
GTTGGCCATG GAACCTGTGC GATAATTGGC TTAATAAAAGT TTAGAGAATT CGCTACGTAA	3900
TAAATAGTAG CAGTAACCAT TGGTGCTAAA ATAAATGGTA TAGCCAAGGC TGGATTATAG	3960
ATAATAGGTA ATCCAAAAAT TAATGGTTCA TTAATATTAA ATAAGGCTGG AACTACAGAT	4020

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GCTCGTCCTA TTGCTTAAG CTGTCAGAT TTAGAGGCAA AAGCAATATA TAAACATAGT	4080
CCTAAAGTTG CACCAGAACC ACCTGCAATT ACAAACATAT TAGAAAATTC ACCTGCAACA	4140
GCGAAGTGCC CGCCAGCAGC ATTTTCAGCC ATGTTAGCAA GAGCAATTGG ACTAACAAAT	4200
GCAAAACAA TGTCGCACC GTGGATACCT ACAATCCAAA GTAGTTGAGT CAATAGATAA	4260
ATAATCATTA AACCAATCCA CGAATTAGTC AGATTGGATA CAAAACAAA TGGAATTGCA	4320
ATGACTTTAA AAATATCTGT TCCCATTGCT ACAAGAAGAC CGTTGATAAA GATAACAAACA	4380
AATGCAACAA CAAATCCCGG AACCAAAGCG GTAAATCCAC GAGAAACTCC TTCTGGAACA	4440
GCTTCAGGCA TTTTAATAAC CCAATTATGT TTAACACACA TACGATAAAAT AAGAACAGTC	4500
ACAATTGCCA TAATGATTGC GGTAAAATC CCTGTTGTCC CAAAACGTGC GACTACATTT	4560
CCCATTGCCC ATCCATCTGC AATTACTGCA CCTTCTTTA GACTTGTAC AGTCTTCATC	4620
ATTCCACCAT CAAAAATGAT TTGCGGTACT GTCATGACAA AAGCCATCAA GGCAAGCAAG	4680
GCACCATTAA GAGGATTCAT ATTGAGTTCT TCTTCCTCTG CATAAATTTC TGTCAATTCA	4740
TATGCAAGTG ATAGAACGAA ATAAAGAGAT AGAGAACCCA TAGTCGCATA GTTGTCAACC	4800
ATGTAAAGTG ATGTGAATTTC ATCAAATGAA GCAGAGAAAA TATCTGCCAC AATTGCCAA	4860
AATGAGAAAG CTTGTGGCAA AATACTGAAT ACCAAAAACA TTGATCCTAC AATAGTAAAT	4920
GGTACAGCAG CCATACCTGC AGCCGTGATA GCACGTACTA CTTTAAACTG AGCAAGTTG	4980
CCCATTGGTC CCATAACATG GTTTCAAGA AAACCAAACA ACCCGTTTG TTGATCCATA	5040
AATAGACCTC CTTAATAAAAA CATAATAATT TTTACTTTCT AAAGACTAGT TTCAAAATACA	5100
AATTATACTA GATCAGGATT ATAAACTAAG TGAGTTCTTT TCCAATTGGA CAAATTGTTG	5160
ATAAGCCTTA TCTGTCGTT TATAAATTTC TTTAATTCTT CTAATGTCTA ACAAACTCAG	5220
AACTAAACCT AATAGAAGAA CTACAAAAC AAATAAACGT GCTACTTGGT TATTTCAAA	5280
AATCGGAAAA AGATTCTTAA ACCAACTGT CCAAGTTAAA ACAAGTAATC CTATTGAAAT	5340
AAGCATTGTT ATTCTAACAA ACATTAGTGT TATTCCCAAC TTTCTTTCC TATTTCCATA	5400
AAGTTTAAAT TGTCACACAG TTGCTAAAAT AGAAAATACT ATGAGCATAA TGGGGAAAAT	5460
AATAATAGGC GAGGGACTAA TAAACTGACT CAAAAGCCAA TAAATATTCC CAAAAAAGAA	5520
GAGTGCTATT GAATAACGTA GAAGAAGATA TCGATTGAAA AAAGTATTAG TTAGAGCCAT	5580
CTCTCGACGT TGTTGTTCAA TCTTTGTCG TTCTTTTTA TCCATATCAT TTCTCCTTA	5640
TATAACAACA CATATTTAGT TAACTTTCTT ATAAAGAGCT AACATTCCT TTGCTACTTC	5700
TAATAATGTC ATAGTGGTCA TTAATGATC TTGAGCATGT ACCATGATAA TTTCAATTTC	5760
AATTTCCACT CCACTTGCGT ATTCTGCAA GAGTTGGTT TGTGCATGAT GCGCTTCAAG	5820

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AATTATCTCA	TTTGATTGAT	TTAATTTACT	TTCTGCATCA	TCAAAACTAC	CTTCTCTCAT	5880
TTTTGCAAAT	GCTTCATGTA	TTTCTGACCT	TGCATTTCCC	GAATGCAGGA	TAATTTCAA	5940
TGCTGCAACC	TGCAGTTCCT	CTTGATTCAT	ATAAACCTCC	TATTTTATCT	TCTCAAATAT	6000
GTAAATAAAA	TCTTCAAAGT	TATTGCAAGA	TATTAGCTGA	TTTGCAATT	CATCATTCTC	6060
TGTCAGAGAG	ACTATCTTT	TAGTCACAGT	TGCCAAACCT	TCGTTCCCAT	ATATTGATGG	6120
AGATAGAAGA	AATACTAGCT	GGACATGTGA	ACTTGATTA	TCCCAGAGTA	ACGAATCTT	6180
ACAAATTGCA	ACCGAAACCT	TTCCCTCTGT	ACCAAAGGGC	TGAATAGGAT	GCGGAACTGC	6240
AATTTTTCA	GAAAAACAA	CTGAACTTAA	TTCTTCGCGC	TGTTTAATTC	CATAAAGTAA	6300
AGATTGTTCA	AACTCATTG	ATTCACCAAC	AGATAAACTC	TCAACCATCT	TTTCAAGTAA	6360
ATTTACCTTG	TCTGATTCA	TACATATTAA	AAAGTTTCT	TTACTAAAAT	ACTGTCTAAA	6420
GCCGTTGTTT	TCAAATTG	TAATCTTG	TGATTGTACA	TAACCTAGAAA	CTTGATCTA	6480
ATCCATAGCT	TTTCTAAATCA	TTTCCATCTC	ATCACTCTTA	AGAAACACAC	TAACTTAAA	6540
AACTGGGATT	TGAAATATA	GATTGATAA	ATCAATAGCT	GACACTATAA	AATCTATTCC	6600
TTTAAGTTTT	TCTTGATTCA	ATTCATAGTA	GCCTATTACA	TCAACAACTT	CTACTCGCTT	6660
CCCAAACCTCC	GTTTCCAAAC	GATTCTTAA	CATTTGGGCT	GCACCAAATC	CTGTTGCACA	6720
AATAGCAAGA	ATATTAAACT	TAGTACTCTC	TTTGCTACGT	TCCATAGCAG	CTAAAAGTG	6780
AAGACTTACA	TATGCTACTT	CATCATCTGA	TATTGTCCAC	TCCAAGAACT	TGTCCATATT	6840
TGCAAGAATT	TCTCTAGTCA	TAAAGAATAT	ATCACTATAA	TTCTGTTAA	TTTCATCTAC	6900
CAAAGGGTTA	TTTAAGGTA	TCCGGCTTTC	AAACGTACT	TGTAGTGTCA	TTAGATGAGT	6960
TATCAATCCT	TCAATTAGTT	GGAAATCTGA	AGAAAAGTTA	TACATATCAT	CTAACCTAA	7020
ATTCTGAAAT	GTTTTAAATA	AAGATTTTT	AAAAACTTCT	TCAGAAATAT	TCTTCTGATT	7080
TTTTTGACAT	TGTTGACTCT	TAGCTAACAA	ATGCAAAGTA	ATGTAGTCTA	TTTCCTGAAC	7140
TGGAAATTCC	TGATTGTTA	CTTCTCTTAC	TTAGAAAGA	ATTCTTTGGG	CAACCTTCT	7200
CTCTATTGCA	TCATCAGTC	TCTGACAGTC	TATATTTTTT	ATTCAAATC	CGGATTTAA	7260
ACGAATCACA	GACAATGCTA	TGTGAACATAC	AAATTCTGT	AGTACAAAT	CAGATAGTT	7320
TAGGTTGGCC	TCTTGGCATT	CATCCAAAAC	AAATTCTAGCA	AATTCTCTA	ATGGAACAGT	7380
TTGATCAAAA	AAGTTAAATT	TTACATAGCA	ATGTATTGTT	TTAAAAAAATT	GATTCTCTAG	7440
GAAATAATT	ATGATAAAAC	GTGTTTATC	ACGTTCCCTCG	CCTGAGACAT	AAACTCCTTT	7500
ATTGCCCTA	CTCTCAATGG	ACAAATTATA	CTCTGATAAC	ATCACTCGTA	TCTTCTGAA	7560

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ATCATGAGAT AATGTTGAAC GACTAACGTA AAGTTCATCA GCTAAATCAT CAAAAAGAAC	7620
TGGAACTTGC TCAAATAATA ATTTATTAA GATAAATACT AAACGATCAT CACCTTTGA	7680
AACCGCAGTT TTCGTATACT CTTCTTCCAG TTCAAGTT TGTCTAACT CCTGGTAAGC	7740
GCCTTGATTC TCACAAAATA TTTGATAACCC TTGACCTTGT TTTGAAATCA ACCGGACTCC	7800
TTGAATAATC ATTGTCTTCT CAATTAATT CAGTACATTA CGGACAGTTC TATCTGAACA	7860
GGATAAATAT TCTGCCAGT CTTTGCTTGT AACAAAACGT TCCTTATTAA TTATTAACAA	7920
TTGAAGGATA TCTTTCTTAA TAATGTTAA CACATTCAATT CCCTCCTAAA ACGTATGTTT	7980
TCATATATTG AAGCATATTA TACACTTAA TCAGTTATA TCAAAACTCAA AACAAATTAT	8040
CTTAACCTAA ATATTTATTG ACATTTCATG TGTTCATCAA ATATTCTCAA GAATCAAATT	8100
AGCCATTTTT TCAATTCCCA TTGGAATAGG AATATAGGCT TGAGGAGGTA TTTGTACAAC	8160
TGGTTTCCCT GCTTTAGAAC CAGCCTCTTC AAATTGCTTA AAGTACATTT TTGTTGAGG	8220
ACTGACAAGA TACAAATCAA AAGCTGCTGC TGCGATAGCT TTCCCTCCTT CAGTAGCACT	8280
AATAGCATCA ACTACAATAT CTTCCCTTT TCCTTTAGA AACTCTGTG TTTTCTGTGC	8340
CATAAGTGAT GAAGACATTC CTGCTGCACA AATAATTAAA GCTTTGCACAA TAATTTTC	8400
TCCTTTCTT AAATCCAATC AAAGCTGTGC TAAGTTGGCT TATTTGTTAT CTATTTTAT	8460
TATAAAATAA AGCGTTCCA ATGACAATTC CCTCATTTC CAAATGATA TGGAAAAAAA	8520
TTATTTATAC TTCAATTAT AAAATAAAAT TATTCCTGAG AGTAGAAATG AAACACTATT	8580
TGCTAAAATC AAAGGCAAGT CTCCTATACG AATACCATGA GCAAGGCCACA ATGCAATACC	8640
AATAACTTGC ATAACATACA TACCTAGAGC AATAGATCCT GTGTCCTTG TCTTAACTAC	8700
ACGAAAAACT TGTGGAAAAA ATGCAAATGT TGTTAAATT GCTGCAATAC TTCCAATCAT	8760
ATGTCACCTC AATATGCTAA ACAAACTGAG AATAATCTCA GTTTGTTAT ACTATTCTAC	8820
TGATTCACCG TTAGATGAAA TAACTCCTT ATACCAGCCA AAAGATTTT TCAGGGAAACG	8880
ATTATAACTT CCCTTCCCAT TATCATCTT ATCTACATAA ATAAAGCCAT AACGTTCCG	8940
CATTTCACCG GTACCAGCTG AAACCAAATC AATACATCCC CATGGAGTAT AACCCATTAA	9000
ATCAACACCA TCTTCAACTA CAGCCTTTT CATTTCACGA ATATGGGCAC CTAGATATTC	9060
AATTCTATAA TCATCATGTA CCATACCATC TGCTGCAACT TGATCTATAG CTCCAAAACC	9120
ATTTTCAACA ATAAAGAGTG GTAAGTGATA GTGGTCTGTA AACCAATTAA ACGCATAACG	9180
CAAACCTTCT GGATCAATTG GCCACTCCCA TTCAGAAGCC TTAACATAAT TATTTTCAC	9240
TAAATCTTCT GTTTCAAGAT AATCAAAATA AGGATTATTT TCACGATGAG AGTCGATAGC	9300
AAAGGACATA TAGTAACTGA AACCAATGTA ATCTACAGTC CCACCAAGTA AATCTTCTTT	9360

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ATCCTGGCA	GTAAAATCAA	CTGAAATACC	TTTCGTTCC	CAATACTTGA	AAATATGCTC	9420
AGGATATTAA	CCTAAAACAT	GCACATCAGC	AAAATAATAA	CGCTTCTGCA	TAGCTTCAT	9480
TGCCATTAAAG	ATATCCTTAG	GATTGCAAGT	AACTGGATAA	ATTGGACACA	TCGCAATCAT	9540
ACAACCTATT	TGAAAATCTG	GATTAATCTC	ATGACCAATT	TTTACAGCTC	GTGCAGAAGC	9600
AACTAATTG	TAATGTGCTG	CTTGATACAT	AATTGCTTCT	CTATTATCAC	CTTCCTCATA	9660
TACAATACCT	GAGTTAGTAA	ATGGTGCAAA	ATCTTCCTGA	TAATTGCTT	GATTATTGAT	9720
TTCATTGAAA	GTCATCCAAT	ATTTAACCTT	ATCTTTGTAA	CGTTTAAATA	CGACTTCTGC	9780
AAAACGAGCA	AAGAAATCAA	TCAATTTCCT	ATTTTCCAA	CCACCATATT	CGGTCACTAA	9840
GTGATAAGGC	ATTTCAAAAT	GAGATAGAGT	GATGACAGGT	TCAATACCAT	TCTTTAAGCA	9900
TTCATCAAAA	AGATTATCAT	AAAACGTAA	TCCTTCTTCA	TTCGGCTCTA	ACTCATCACC	9960
TTTGAAAG	ATACGGTCC	ATGCAATAGA	GGTACGGAAG	CACTTGAATC	CCATTTCAGC	10020
AAAAAGTGC	ATATCTTCTT	TATAACGGTG	ATAAAAATCT	ATCGCCTCAT	GATTTGGATA	10080
ATATTTACCC	TCTAAAACTC	CCAAAGTAAT	TTCACGAGCT	ACTCCATGAC	GACCAGCAGT	10140
CATAACATCA	GCAACACTAA	TTCCCTTGCC	ACCTTCTTGC	CATCCACCTT	CAAGTTGATG	10200
AGCAGCAACA	GCACCACCCC	ATAAAAATCC	ATCTTTAAA	GTAGTCATCT	TTTTCCCTCC	10260
TGACTTTGAT	ACTCTTATTA	TAAACCTTAA	ACCAAAAGAT	AAAAACGCAT	TCTTTTCCCT	10320
TATTGTTAAG	GAAAGAAGTA	ATTTTAATG	GAAATAGAAC	AATATCTTCT	TGTATTCTCG	10380
TAATGATATC	TTTACGATTT	TCAATACTTT	CAAAC TACAA	AAACTCTCAC	AATAATTCTA	10440
ATTCCTGTG	TCTATAAACG	ACTTATCGCT	TTCTGGCATC	CCAGAACATC	CTTCTATATA	10500
ACGTTCAACT	TGCATCTGCA	AGTGATATTT	TTTTCTTAAA	TCTAAGATT	TCTGCATTGT	10560
CTTGATGTA	TAATGTTAT	CTAAAGTTTC	TTGATTTATC	CACTGATCAA	TAAGGAGAAT	10620
AGTCCCTCT	TTTCAATTG	GTAAAAAATA	TTCGTATTTTC	AAGTACCTT	TTGATTTCT	10680
AATTTCTTA	ACAAGGCCAC	TATCAAGCAT	TTCTCTTGCA	AACTTTATTG	CACTATCTCC	10740
ATCACCTTA	TAATATACAT	GAATAGTCAA	TGTCATCTTA	TATCCTCCAA	AATCATCCTT	10800
CAATTAAA	AAAACAAGTT	TAGATGAGGA	TCTAAACTTG	TTTTTATGA	ACTAATTATC	10860
TAACGTTTCG	CCATTACTTT	CAATCACTTC	TTTATACCAA	AAAATGATT	TTTTCTTATA	10920
GCGATTATA	GTCAATTGAA	ACAAGAGCAG	GACAAAAGAG	CCTCATAAAA	GGTATTGCAA	10980
CTTGGTAATA	CCTTTTGAG	GTGCTTTTG	ATATGAGCCC	ATGTTTCTC	AATAGGATTG	11040
TACTCAGGTG	AGTAGGGAGG	AAGAGGTAAA	AGTTTATACC	CAAACCTTC	ACACAAGAGT	11100

838	
TCTAGCTTCC CCATTCATG GAATCTTGCA TTATCCATAA TAATAACCGA TGGTGTGGTT	11160
AATGTTGGTA AGAGAAACTT CTGAAACCAA GCTTCAAAAA AGTCGCTCGT CATCGTCTCT	11220
TCGTAAGTCA TTGGAGCGAT TAACTCACCA TTTGTTAGAC CTGCAACCAA AGAAATCCTC	11280
TGATATCTTC TTCCAGATAC TTT	11303

(2) INFORMATION FOR SEQ ID NO: 116:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 3112 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 116:

CCTTAGAGATT CCACCTGCCA GAGGAATTGA TTGCCAAAC GCCCCTGAA AAACGTGATG	60
CCTCCAAACT CCTCATCGTC AACCGTGAGA CAGGAGAAAT GCAAGATAAA CATTTCCACT	120
CTATTATTGA TATGCTGGAA CCTGGTGATG CCCTTGTCAT GAACGACACC CGAGTTCTCC	180
CTGCCCGCCT CTATGGTCAA AAAGTGGAGA CAGGAGGTCA TGTGGAACCTT CTCCCTCTTA	240
AGAACACTAG TGGAGACGAG TGGGAAGTTC TGGCTAAACC TGCCAAACGC CTCAAGGTGCG	300
GTACTCGTAT CAGCTTGTT GATGGCCGCC TCAGCGCTGT CGTTACAGAA GAATTGACCC	360
ACGGGGGACG CATTGTCGC CTTGAATACC AAGGAATTTC CCTAGAAGTC TTGGAAAGTC	420
TGGGAGAAAT GCCTCTGCCA CCTTATATCC ACGAAAATT AGATGACCGT GAACGTTATC	480
AAACCGTCTA CGCCAAGGAA AGTGGCTCTG CTGCAGCACCC GACTGCTGGT CTTCACTTCA	540
CCAAAGAACT GCTGGCAGAA ATCCAAGCTA AGGGTGTCA TCTAGTCTAT CTGACTCTCC	600
ATGTCGGACT CGGAACCTTT AGACCTGTTT CTGTGGATAA TCTGGACGAA CACGAAATGC	660
ACTCAGAGTT CTATCAAATT TCTGAGGAAG CTGCTGCCAC CCTTCGCTCT GTCAAAAAAA	720
ATGGTGGTCG TGTCATCGCT GTCGGAACCA CTTCTATCCG CACCTTGGAA ACTATTGGTT	780
CCAAGTTGTA TGGGAAATC CAAGCAGATT CTGGTTGGAC CAATATCTT ATCAAACCTG	840
GGTATGAGTG GAAGGTCGTG GATGCCCTCT CAACCAACTT CCACCTGCCA AAATCAACTC	900
TGGTCATGTT GGTTCTGCC TTTGCAGGCC GTGAATTAGT CTTAGATGCC TACCACCATT	960
CCATCCAAGA ACACCTACCGC TTCTTCAGTT TTGGTGACGC CATGTTTATT TATTGAGAAA	1020
GAATTCTCT AAATCTCTA ATACCAATAA ATCGCTAAGA TATTATTCAGA AAGAACATCT	1080
ACAATTGAAA CTCTAGCTAG CTGTAGAAGA GGCCTAGTAC ATTGAAATTA AAATGCTTCC	1140
CCCTAGCTTC GAAAATATTG CCATAGATTG CGTTGACTCT CCAAATTGAT TCATCTATAT	1200

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TTTATTCAG CTTCCCTATAC	TTTCTTCGCT GTTTGAAAT	CAAATGCAA GACACATGAG	1260
TAGCACCATA TTTGTTACTC	TTATCTGTCC TCTCAAGAGA	CTATTATGAG TTATTCAGA	1320
ATCATTCACT ACTTTGACCC	TGACTCTCCT TAGTCTCAA	ATCAAAGACT TATACTCTC	1380
AAAAATCTCT	TCAAACCGCG TCAACGTCAC	CTTGGATTAT ATATGTGatC	TGaCTTCGTC
AGTTCTATCT	ACAACCTCAA AGCAGTACTT	TGAGCAACCT GCGACTAGTT	1440
CTCTTGATT TTCATTGAGT	ATTAAACAAA AAGTGAACAA	ATCTGAATTC TAATGTACAG	1500
AAGACTAGGC TTGTTCACTT	TTTTATAGTC GCTATAAGAT	GACCTTATCT ATAGCTTTT	1560
ATATATAATT ATATATTCA	ACATACTATT ATCAATTTC	TGCGCAGGGAG GAATCTGTTA	1620
ACGCACCCAT TCACCATTAT	CATTGACTCT ATAGCCATCT	ATACTTGTAT TGACCGCTAA	1680
CTCACCCGAT GTATTACAT	AATACCATT ACCACCAACT	TGGAACCATT GATTGACTTT	1740
CATAGAACCG TTGCTGTTGA	GGTAGTACCA TGAECTATTA	ACTTGTACCC AACCTGTTGC	1800
CATGGAACCA TCAGTATTAT	AAAAATACCA CATAACCATT	TCTTGTTCAG AGTCTGTTGT	1860
TGGAGCAACT GCTTAGCTG	GTTCTACTGC TACATCTGTT	CCTTGGTTAG ATGTAACAGA	1920
TACAGGATAC GAAGGAATAG	ATGATTGCTC AGGAACAACA	ACTTTTCAG GTTCTCTCGT	1980
CCCTCTCCTT ATACGTCTT	TTACCATCTC TTTAGTAATT	TGACGAGAAAG TAGTTCTTC	2040
AATTGTTCCA TCACGTTCAT	CTACAGTATA GATTGTAGTA	AGAGTAATTT ACCAATTCT	2100
CCTACTCTT CTACTTCTTG	ACTTTTATCA AGAGTTGGCC	CATCGAGATA TTCTGTTCG	2160
ATTGGAATTT CTTGGACAAG	AACTTGGGGC TTGGTTCTTT	TTTTAACAAAC TCTTGTTCGA	2220
GAGTCTTTT TTTGACTTAA	AGTACTCTCA GTTACTTGTC	CACTCTTCC ATCTACATTA	2280
TAAGTTATCG TTGTAACTGT	TTTCCCATTCA TTTCTAGAG	TAATCTCTTG CTCTGTCCT	2340
GCAGAAAGGT CATTGCTG	TTCATATTAA GTAGCAAATG	GAACAAGAAC TTCTTCAACC	2400
TTGCTTTAG CTGAACTTT	GATAACTGTA TCCGTGGCTT	CTTTTCTATC AACAGTAACC	2460
TGTCGGTAA CATAACCAGT	CTCTGGATTA ACATCGTAGG	TCCTTGTGCGT AGTTACATAG	2520
CCATCCTCTC CATCAATTGT	AACAGGATTTC TCACTACGGT	CTTTTGTTC ATCTTTTCA	2580
TAACGAATTC GCGTACTTGA	AATTTCTTG GTTACTACCT	TAGGTTTAGT CGCTACTTTT	2640
ACAATAATAT CCCCATTGTC	AGCGTCATCA TACTCTATTC	CCTCTCTTTT ATCTCTAGTA	2700
TCATCTCTGA CATATTGAAT	CCCATCAGCA GCATGAACAA	AACTTGTATT CAGATTCTC	2760
CTAAAAATAA AGTAGCCCCG	ATTACCGCAG AACCAAAAT	CTTCCGAGT TTACGTATTG	2820
CATAGCGCTT ATTAGTATTA	GATTTGCCA TTACATCCTA	CTTCTAGTAT AGCATCTTT	2880
			2940

840		
CTATCAAACG TTAAACAATA TACGTTATAT ATAAAATAGA CTTAGAACATGA TATATTGATT		3000
ATTGAACTAA CACTTTAAT ATATCGTAAT CAATCTCATA TATAAAGGAT TGCAAGACATC		3060
TTATCTAAAT ACATGCGAAT ATATTTAGAT ACAAACATTC CAACTTGATA AT		3112

(2) INFORMATION FOR SEQ ID NO: 117:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 4327 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 117:

CCCCAAAATC	TCTTCAAACC	ACGTCAGCTT	CGCCTTGCCG	TAGTATGGTT	ACTGACTTCG	60
TCAGTTCTAT	CCACAACCTC	AAAACAGTGT	TTTGAGCATC	ATGCgGCTAG	CTTCTTAGTT	120
TGCTCTTGGA	TTTTCATTGA	GTATAAAAAC	AGATGAGTTT	CTGTTTTCTT	TTTATGGACT	180
ATAAAATGTC	AGCTGAAACT	ACTTTCAAGG	ACATTATTAT	ATAAAAGAAC	TTTTTGAAAC	240
TAAAATCTAC	TATATTACAC	TATATTGAAA	GCGTTTTAAA	AATGAGGTAT	AATAAATTAA	300
CTAACGCTTA	AAAAAAGTGA	TAGAATCTAT	TTTTATGTAT	ATTTAAAGAT	AGATTGCTGT	360
AAAAATAGTA	GTAGCTATGC	GAAATAACAG	ATAGAGAGAA	GGGATTGAAG	CTTAGAAAAG	420
GGGAATAATA	TGATATTTAA	GGCATTCAAG	ACAAAAAAAGC	AGAGAAAAAG	ACAAGTTGAA	480
CTACTTTGGA	CAGTTTTTT	CGACAGTTTT	CTGATTGATT	TATTCTTCA	CTTATTTGGG	540
ATTGTCCCCT	TTAAGCTGGA	TAAGATTCTG	ATTGTGAGCT	TGATTATATT	TCCCATTATT	600
TCTACAAGTA	TTTATGCTTA	TGAAAAGCTA	TTTGAAAAAG	TGTCGATAA	GGATTGAGCA	660
GGAAGTATGG	TGTAATAGC	ATAGGCTGAT	GTCCCATCATT	TGCTTATAAA	GAGATATTTT	720
AGTTAATTG	CAGCGGTGTC	CTGGTAGATA	AACTAGATTG	GCAGGAGTCT	GATTGGAGAA	780
AGGAGAGGGG	AAAATGGCA	CCAATTGAG	ATAGTTGTT	TAGTCATTT	TTGTCATTAA	840
AATGAACGT	AGTAAAAGAA	AGTTAATAAA	AGACAAACTA	AGTGCATTT	CTGGAGTAAA	900
TGTCTTATTT	CAGAAATCGG	GATATAGATA	TAGAGAGGAT	CAGTATGAAT	CGGAGTGTTC	960
AAGAACGTTAA	GTGTCGTTAT	AGCATTAGGA	AACTATCGGT	AGGAGCGGTT	TCTATGATTG	1020
TAGGAGCAGT	GGTATTGGA	ACGTCTCCTG	TTTAGCTCA	AGAAGGGCA	AGTGAGCAAC	1080
CTCTGGCAAA	TGAAACTCAA	CTTCGGGGG	AGAGCTCAAC	CCTAACTGAT	ACAGAAAAGA	1140
GCCAGCCTTC	TTCAGAGACT	GAACCTTCTG	GCAATAAGCA	AGAACAAAGAA	AGGAAAGATA	1200
AGCAAGAAGA	AAAAATTCCA	AGAGATTACT	ATGCACGAGA	TTTGGAAAAT	GTGCGAACAG	1260

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TGATAGAAAA AGAAGATGTT GAAACCAATG CTTCAAATGG TCAGAGAGTT GATTATCAA	1320
GTGAACTAGA TAAACTAAAG AAACTTGAAA ACGCAACAGT TCACATGGAG TTTAACCCAG	1380
ATGCCAAGGC CCCAGCATTC TATAATCTCT TTTCTGTGTC AAGTGCTACT AAAAAGATG	1440
AGTACTTCAC TATGGCAGTT TACAATAATA CTGCTACTCT AGAGGGCGT GGTCGGATG	1500
GGAAACAGTT TTACAATAAT TACAACGATG CACCCTAAA AGTTAACCA GGTCAGTGGA	1560
ATTCTGTGAC TTTCACAGTT GAAAACCGA CAGCAGAACT ACCTAAAGGC CGAGTCCGCC	1620
TCTACGTAAA CGGGTATTA TCTCGAACAA GTCTGAGATC TGGCAATTTC ATTAAAGATA	1680
TGCCAGATGT AACGCATGTG CAAATCGGAG CAACCAAGCG TGCCAACAAT ACGTTTGGG	1740
GGTCAAATCT ACAGATTGG AATCTCACTG TGTATAATCG TGCTTAACA CCAGAAGAGG	1800
TACAAAAACG TAGTCAACTT TTTAACGCT CAGATTAGA AAAAAGACTA CCTGAAGGAG	1860
CGGCTTTAAC AGAGAAAACG GACATATTG AAAGCGGGCG TAACGGTAAC CCAAATAAAG	1920
ATGGAATCAA GAGTTATCGT ATTCCAGCAC TTCTCAAGAC AGATAAAGGA ACTTTGATCG	1980
CAGGTGCAGA TGAACGCCGT CTCCATTGCA GTGACTGGGG TGATATCGGT ATGGTCATCA	2040
GACGTAGTGA AGATAATGGT AAAACTGGG GTGACCGAGT AACCATTACC AACTTACGTG	2100
ACAATCCAAA AGCTTCTGAC CCATCGATCG GTTCACCAGT GAATATCGAT ATGGTGTGG	2160
TTCAAGATCC TGAAACCAA CGAATCTTT CTATCTATGA CATGTTCCA GAAGGGAAGG	2220
GAATCTTGG AATGTCTTCA CAAAAAGAAG AAGCCTACAA AAAAATCGAT GGAAAACCT	2280
ATCAAATCCT CTACCGTGAA GGAGAAAAGG GAGCTTATAC CATTGAGAA AATGGTACTG	2340
TCTATACACC AGATGGTAAG GCGACAGACT ATCGCGTTGT TGTAGATCCT GTTAAACCA	2400
CCTATAGCGA CAAGGGTGAT CTATACAAGG GTGACCAATT ACTAGGAAAT ATCTACTTCA	2460
CAACAAACAA AACTTCTCCA TTTAGAATTG CCAAGGATAG CTATCTATGG ATGTCTACA	2520
GTGATGACGA CGGGAAAGACA TGGTCAGCTC CTCAAGATAT TACTCCGATG GTCAAAGCCG	2580
ATTGGATGAA ATTCTTGGGT GTAGGTCTG GAACAGGAAT TGTACTTCGG AATGGGCCTC	2640
ACAAGGGACG GATTTGATA CCGGTTTATA CGACTAATAA TGTATCTCAC TTAGATGGCT	2700
CGCAATCTTC TCGTGTCAATC TATTCAAGATG ATCATGGAAA AACTTGGCAT GCTGGAGAAG	2760
CGGTCAACGA TAACCGTCAG GTAGACGGTC AAAAGATCCA CTCTCTACG ATGAACAATA	2820
GACGTGCGCA AAATACAGAA TCAACGGTGG TACAACAAA CAATGGAGAT GTTAAACTCT	2880
TTATGCGTGG TTTGACTGGA GATCTTCAGG TTGCTACAAG TAAAGACCGA GGAGTGACTT	2940
GGGAGAAGGA TATCAAACGT TATCCACAGG TTAAAGATGT CTATGTTCAA ATGTCTGCTA	3000

842	
TCCATACGAT GCACGAAGGA AAAGAATACA TCATCCTCAG TAATGCAGGT GGACCGAAC	3060
GTGAAAATGG GATGGTCCAC TTGGCACGTG TCGAAGAAAA TGGTGAGTTG ACTTGGCTCA	3120
AACACAATCC AATTCAAAAAA GGAGAGTTTG CCTATAATTC GCTCCAAGAA TTAGGAAATG	3180
GGGAGTATGG CATCTGTAT GAACATACTG AAAAGGACA AAATGCCTAT ACCCTATCAT	3240
TTAGAAAATT TAATTGGAA TTTTGAGCA AAAATCTGAT TTCTCCTACC GAAGCGAACT	3300
AGAGAGATGG GCAAAGGAGA GATGGGCAA GGAGTTATTG GCTTGGAGTT CGACTCAGAA	3360
GTATTGGTCA ACAAGGCTCC AACCCCTCAA TTGGCAAATG GTAAAACAGC GACTTCCCTA	3420
ACCCAGTATG ATAGCAAGAC CTTGTTGTTT GCAGTAGATA AGGAAGATAT CGGACAGGAA	3480
ATTATTGGTA TAGCTAAAGG AACCATCGAA AGTATGCATA ATCTTCCTGT AAATCTAGCA	3540
GGTGCCAGAG TTCCTGGCGG AGTAAATGGT AGCAAAGCAG CGGTGCATGA AGTTCCAGAA	3600
TTTACAGGGG GAGTTAATGG TACAGAGCCA GCTGTTCATG AAATCGCAGA GTATAAGGAA	3660
TCTGATTCGC TTGTAACCTCT TACTACAAAA AAAGATTATA CTTACAAAGC TCCTCTTGCT	3720
CAGCAGGCAC TTCCTGAAAC AGGAAACAAG GAGAGTGACC TCCTAGCTTC ACTAGGACTA	3780
ACAGCTTCTC TCCTTGGTCT GTTACGCTA GGGAAAAAGA GAGAACATA AGAGAAGAAT	3840
TCTAAACATT TGATTTGTA AAAATGGCTC TTTGTCAACT GTAGTGGTT GAAGTCAGCT	3900
AAGCTCGAGA AAGGACAAAT TTTGTCTTT CTTTTTGAT ATTCAAGAGCG ATAAAAATCC	3960
GTTTTTGAA GTTTCAAAG TTCCGAAAC CAAAGGCATT GCGCTTGATA AGTTGATGA	4020
GATTATTGGT CGCTTCCAAT TTGGCGTTAG AATAGTGTAG TTGAAGGGCG TTGACGATTT	4080
TCTCTTGTC CTTTAGAAAG GTTTAAAGA CAGTCTGAAA AAGAGGATGA ACCTGCTTA	4140
GATTGTCCTC AATGAGTCCG AAAAATTCT CCGGTTCCCTT ATTCTGAAAG TGAAACAGCA	4200
AGAGTTGATA GAGCTGATAG TGATGTTCA AGTCTTGTGA ATAGCTAAA AGCTTGTAA	4260
AAATCTTTT ATTGGTTAAA TGCATACGAA AAGTAGGGCG ATAAAAATGT TTATCGCTGA	4320
GTTCACG	4327

(2) INFORMATION FOR SEQ ID NO: 118:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 3521 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 118:

CTCTGGCCCT GCCACTCCAA CGTTTGTC GGGTGCTTT TTCATAAAGG AGTTCTTATG

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TTAGATATCA AACGTATTG TACAGATTT GAAGCTGTCG CAGAAAAATT AGCTACACGT	120
GGTGTAGATG CTGCTGCTT GAATGAAATG AAAGAAATCG ATGCTAACAG TCGTAACATC	180
TTGGTCAAGG TTGAAACTCT CAAAGCAGAA CGTAACACAG TTTCTGCTGA GATTGCCAA	240
GCTAAGCGCA ACAAGGAAAA TACAGATGAC AAGATTGCTG CCATGCAAAA TCTATCTGCT	300
GAGGTTAAAG CCTTGGATGC TGAATTGGCA GAAATCGATG CTAATTGAC AGAATTTACA	360
ACGACTCTTC CAAATATCCC AGCTGACAGC GTTCCCTGTTG GGGCTGACGA AGACCGACAAT	420
GTGGAAGTTC GCCGTTGGGG TACTCCACGC GAGTTTGACT TCGAACCTAA AGCTCACTGG	480
GATCTCGGTG AAGACCTTGG TATCCTTGAC TGGGAACGCG GTGGTAAGGT AACAGGCGCT	540
CGCTTCCTCT TCTATAAAGG CCTCGGTGCT CGTTTGGAAC GTGCTATCTA CAACTTTATG	600
TTGGATGAAC ATGGAAAAGA AGGCTATACT GAAGTCATCA CACCTTACAT AGTCAACCAT	660
GATTCTATGT TTGGTACTGG TCAGTATCCA AAATTAAAGG AAGATACTTT TGAACTCAGC	720
GATACCAACT TTGTCTTGAT TCCAAGTGC GAAGTTCCCTC TGACAAACTA CTACCGTGAT	780
GAAATCTTAG ACGGCAAAGA TCTTCCAATC TACTTCACTG CCATGAGTCC GTCATTCCGT	840
TCTGAGGCTG GTTCTGCCGG TCGTGATACG CGTGGCTTGA TCCGTTGCA CCAATTCCAC	900
AAGGTTGAAA TGGTCAAATT TGCCAAACCA GAAGAATCTT ACGAAGAATT GGAAAAAATG	960
ACAGCCAACG CTGAAAACAT TCTTCAAAAA CTCAACCTTC CATAACCGTGT CGTTGCTCTC	1020
TCTACTGGAG ATATGGCCTT CTCAGCTGCG AAGACTTACG ACTTGGAAAGT GTGGATTCCA	1080
GCACAAAACA ATTACCGTGA AATCTCAAGC TGTCAAACA CAGAAGATTT CCAAGCCCGT	1140
CGTGCCAAA TCCGTTACCG TGATGAAGCA GATGGCAAGG TGAAACCTCT TCATAACCTTG	1200
AACGGTTCTG GACTTGCAGT TGGACGTACA GTGGCTGCA TTCTTGAAAA TTACCAAAAT	1260
GAAGATGGTT CTGTGACCAT CCCAGAAGCA CTTCGTCCAT ACATGGGTGG AGCTGAAGTC	1320
ATCAAACCAT AAAAATAAG GTTAGCTAT TTCTAGCTAG ACCTTTTTTC GTAACCAAAT	1380
CAGATAAGCA CCTAGTACAA AGAATAAAAT AGTTAGGCAT ATAATGGTTT CAGCCAATAC	1440
CAGGTAATCC AGAAATGGAA GTTCAAAAT TCCCTGAGCC ATCTTGAGCG AGGTCGCTGT	1500
GATAATGGTT GGGAAAGGTGA GGGCTGAGAA GGCTGGTTGA AAACCTTGTGTT TTAAATGTT	1560
GGGCAGACGA GTTAAACAA AGAAAAAGAA GGATTGAGAA GCCAAATCA TGACAATCAA	1620
GACCCAAGTC GGCAGGCTGG TTCCCTCTAC TCGAACTAGA GAAGCCAAGA GTAGAGAGAA	1680
AGGAGCACAG TAGATTCTT CTTGTCCAAG CAAGGCTAGT GGGAGTGGAT GTTTCTTAA	1740
ATCGCTATAA ATAAGGGGAT AGAGATAGAA GGTCAAGAGA AAACCAAAAC TCAAGGTCGC	1800

844	
ATAGGCAATT TCGATAATAC CTACCAGAGG ATAGGTCAAG GCAGCCACTG CTATCCCCAC	1860
ATAGAGAACC GTCCAGCTTG GAGTGGCATG AACCCCTCCGC CCTGGACAAG CAAACTTGAT	1920
GGTAAAACCA GCAATCAAGG TCAAATCCAA GAGAAATGAA AACCACCAAA TCCCTTGTGC	1980
TACCAAAGGA AGATAAGAGA ATACCGAAA GACATAGTC GATAAAATCA TCCCAGCCAT	2040
AGGAAAGGTT GCCATTCCTG ACAAAAGAGG GGGCTTGGTC AATTCTTGCT TGTTTCTTT	2100
CCAATTAAAG AGATGCAGAA TTAGAAAGTA AATCCATAAA ACCAAACCAA TCAGACTAAA	2160
AAGATGGGAT AGAACCGGCA ACGTATCTAA AATAAGATT CCAGCTCCTG CCAAACCTAG	2220
CAAACAACCT GAAAATACTA AGGGGAGTTT TTTCATCCTA ACCTCCAATA ATCATGTTAG	2280
TTTCAGTATA ACATAAAAGC GCTTAAATGA GGATTTAAAA AAACGAGTCC GCTTATTTC	2340
GACTTCATTT TACTCAGATA TGAATTAGGC ATAAGGTTGC AATTCTGGAT TAATTGGTGT	2400
ATTAGCTAAG TTGTTGGCAT AGTTACAGAG GATTGCTAGG CTGACACCAA AAACCACATC	2460
CAAGGCATTT TGTTGAGTGT AGCCAGCTTC TAAAAACTCA GACAAGGCTT CATCTCCTAC	2520
ACGACCCCTG GTATTGATAA CTGCCAAGGT AAACCTAGCT AGGGTATCCA ATTAGGATC	2580
TGTTTCAATT GGAGTACGAT TCGGAAGAGC TTGAATCAAG TCATCATICA TCTGGATTG	2640
TTTGATGGAA AAGGCTGTGT GACCTGCGAC ACAGAAGGCA CAACCATTGG TCACGGCTGC	2700
CGTGATTTCG ACCACTTCAC GCTCAACGGG TGTCAGGCTG TTGCGACGGT GGATAGATGA	2760
GACAATTGG TAGGCTCTA AAAACAGTCGG GGCATTGGCC AAGAGACCGA TTAGGTTGGG	2820
AATATAGCCA TTGTTGTCTT TTTCTACTGT TTCAAGAATT TCTTTCACTT CTGCTGGTGC	2880
TGACTCTACT GTATGGATAG TAAATGTTGT CATAAGATAC CTCTTTCTT ATTATTGACA	2940
CTAATATTAT TGGAAAATCT TATAAAATCC TGATTCTAA GTTTATCTAA GATAAGCTT	3000
TATTCTCTCA TAAGATTTTC GTTGTATAT TAGTTTATCA CACTTCCAAT CACTTGTATA	3060
ATATATATTA TATATCAGGC TGATAAAAAT TATTATAGG CAAAAAAATC ACACGAGCTG	3120
TGTGATTCCA TTATTTGTCA AAATACTTTT TAGTTTCAGC AATAACGACT GGCGACAAGA	3180
CCAAGAGGGC AATCAAGTTT GGCAGAGCCA TCAAGGCCTT AACGATATCT GCGATAATCC	3240
AGACCATATC CAACTCGATA AATCCTCCTA ACAAGACCAT GAGCACAAAA ACCACACGGT	3300
AGAGCCAGAT AAAGCGAACCC CAAAGAGGA ACTCAAAACA GCGTTCTCCG TAATAGTTCC	3360
AACCTAGAAT CGTTGAAAG GCAAAAGTA CAAGGAAGAT GGTCAAGAGA GCAGGCCAA	3420
AGTGTGAAAA GTTTGTTGAG AAAGCTGACT GAGTCAAGGC AACCCCATTC AAGTCACCGC	3480
TCCAAACTCC AGTTACCAAG ATGGTCAAAC CAGTTAGAGT A	3521

(2) INFORMATION FOR SEQ ID NO: 119:

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(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1968 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 119:

AACCTGGGCA	AGCAAGCTAA	AAGCAATGGG	ACCTGGAATC	CTAATGGCAA	CTGCCGCTGT	60
TGGAGGTTCC	CACATTGTAT	CCTCAACTCA	AGCTGGCGGT	TCTTACGGTT	GGTCTCTACT	120
TCTCTTGGTC	ATCTTAGCCA	ATGTCTTTAA	ATATCCATT	TTCCGTTTG	GTGCTGAATA	180
CACAGCTGAT	ACTGGAAAGA	CTTGGTTGA	AGGTTATGCC	GAAAAAGGAA	AACTCTATCT	240
CTGGATTTTC	TTTATCCTCA	ATGTCTTTTC	GGCTATGGTC	AACACGGCTG	GTGTTGCCAT	300
TCTGTGCTCA	GCTATCATCG	CCAGTGCCCT	CCCAATGATT	GGACTTAGCA	TTACTCAGTG	360
GTCCTCTATT	CTCGTTGCAA	TCATTTGGGC	TATGCTACTC	TTTGGAGGCT	ACAAACTTTT	420
AGACGGCATG	GTCAAATGGA	TTATGTCTGC	CTTAACCATT	GCGACTGTTC	TTGCAGTTAT	480
CATTGCGGCG	GTCAAGCATC	CAGAACATACAG	TTCTGATTTT	GTCGAGAAGA	CACCTGGCA	540
AATGGCAGCT	CTGCCCTTCA	TCGTCTCCCT	CCTAGGATGG	ATGCCGGCTC	CTATTGAAAT	600
TTCAGCCATC	AATTCACTTT	GGTCAGCTGA	AAAGAGAAAAG	ACCGTCAACT	TTAACACAGA	660
AGACGCTCTG	TTTGACTTTA	ACACTGGTTA	TATTGGAACA	GCTATCCTAG	CCGCTTCTTT	720
TGTGGCACTG	GGAGCACTGA	TTCAGTATCC	TACAGGGCAG	GCGGTTGAAG	CTGCTTCAGC	780
CAAATACATC	TCTCAATTG	TGGGCATGTA	TGCCTCTGTT	CTTGGCGAAT	GGTCCCGTTA	840
CTTGATTACC	TTTATTGCCT	TCCTCTGTAT	CTTTGGAACA	GTTATAACTG	TTATCGATGG	900
CTATTCTCGC	GTAAATCAGG	AATCTCTCCG	ACTGCTAATC	AGTCAAAAG	AGGACAATCG	960
TAAATCTTG	AACATCTGGA	TGACCATCAC	TGCTATCATC	GGTATCGTCA	TTATCAAGTT	1020
CTTCGCTGGT	CAGGTTCAA	CCATGCTCCG	CTTGCCATG	ATTGGCTCTT	TCCTGACAAC	1080
ACCTTTCTTT	GCTCTTTGA	ATTACGCCCTT	GGTAACCGGT	GAAAACAAAA	ATCTTCCTTC	1140
TTGGCTCAA	CACCTTGCA	TTGCGGGATT	GATTTCCCTC	TTTGCTTCGC	CATCTCTTT	1200
ATCTACGCAC	TCGCAATCGG	AAAACCAGGG	TAAGGGACAA	GCGCGAGATG	AAGATAAGGT	1260
TTCATTCAA	GAGAAAATTC	AGCAAATATT	TCTATGATAA	AAAGCATAAG	AAACAGGTTT	1320
TGAAGACCTG	AACTTATGCT	TTTTTACGTT	CTTAAAGACT	GTTTATACTC	AAAAAACAGT	1380
TGAACAACTT	CAACCACCTC	TTATAAGAAC	TTTATACTAT	TCGAGAATCT	CTTCAAACCCA	1440

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CGTCAGCTCT ATCTGCAACC TCAAAGCTGT GCTTGAGCA ACCTGCGACT AGCTTCCTAG	1500
TTTGCTCTT GATTTTCATT GAGTATTAAT TCTCCTTTTC CAACTCATAC AAATCTGCAGA	1560
TAATAGCTGC GACATGTTG ATATCTTCCA GCATGCCTCG CATTCAAAG TCAGCCAATA	1620
CAGGGAAGCC AAAGCGTTGA CTGTATTGCT TGGCTGTTAG GCAGTATTGG TTATTAAGT	1680
TACGATTTC TGACCCAACC ACACCAAAAC ACTTACTAGC ATTGTTACCA TAGGCAATAA	1740
AATCTCCAC CGGTGTCGTC AAAATCTCAA CATCTCCGTAT ATCCACGCCA TTCCCACCTT	1800
CGAGATAGGT CGGCAAAAAA GCGACATAGG GATGGTCCAT TTCATAGAAA TTTTGCCCTT	1860
CCTTGACCAA ATCCTTGATA TGAATCTTTT GAACCTCAAT CCCTTTGTAC TGGGACAAGA	1920
GATAGTCTTT CAAGCGCGTC ACAAAACTTT CAGTGTGCCC ACTCAAGG	1968

(2) INFORMATION FOR SEQ ID NO: 120:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7172 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 120:

CCGCATTTT TATCACTAGA CTCGAGACAT CTTTGAGTG GCTCTGCTC TCTGGTTAA	60
TTTTCTTCCT TGCTCAAGGA CTCCTGCTAT TTCTCTGGT CGTCCGACTC AAACATCAAT	120
TCGCTGAGAT TTATCCTCAA ATCAATAAAA AGATTCGTTT CTACTATTAA GGGGTTCTCA	180
CCATTGATT TCTATTTTTT GTTCTCTTAG CCTTCATTAG TTCTCAGCGT TTTTCATCTC	240
TTATGCCAAT CATCACTGCT TGCCATTCTA CTTTTTATTA TATGACAGCT GACTACCTAA	300
GAGAAAACCA TCCAGACTTT TACGACAAAC ACATCTTTT ATGGGAGTGT CTCTAAAGAA	360
AAGGAGGTTT TAGCATGAAA AAAATCATCT TCATCAAAAC CATTCAACTC CTTGTCATTG	420
ATGGAATCAT GCTGGCATT TTGACATTAA AAAGGGGGCT TACTTGGAC TGGATTTGA	480
TTTATAGCGG TTGGCTCATT TTCTTCATC CTGTGCTATT GACCTATCTT TCAAACCAAC	540
TTTGTGACCA CTTTAGTTAA CTCTATTCCC AGATTAGACC GAGATTCTGG CGTTTGCTT	600
TACAAATTCT CCTATGGGAT AGCCTGATGA TTCTCTCCTT GGTGTCTTTA AGTGATATTCA	660
CACTTTCCCT TCAGGGAACCT CTCCTCATCC TAGGACATCT CATCCCTTCC TATCGCATCT	720
GCCAAAGCCT GAAAAGAGAC TTCCCCAAG CATATCAAGA ACCGATTCTT TTTTGGAGTA	780
TTTTATGATA GATGAGAAAG ACCAAGCCGA CTGGGCTTGG TCTTTCTTAT CTCTTTTAG	840
TATCTAGGAT AATGGTAACA GGTCCATTAT TAACCAGCTC AACCTGCATA TCTGCTCCAA	900

847

AGATGCCTGT	CTGAACGGGC	ACTTCTTGC	GCTAATTTTG	ATTGAAAGCA	TCATAGAAGT	960
CTGATGCCAT	ATCAGGTTA	GCTGCCCTG	TAAAGgCTGG	ACGATTGCCT	CTCTTAGTAT	1020
CCGCAAAGAG	GGTAAACTGA	GAAATAGAGA	GGATTCTCC	TTCAATATCT	TTGACAGACA	1080
GGTTCATCTT	GCCTCTGCG	TCTGAAAAAA	TCCGCATATT	GACCAGTTT	CTCACAGCAT	1140
AGTCCAATC	TTCCTCTGG	TCCTCTGGTC	CAACACCAAC	CAGCAATAAA	AGTCCCTGAT	1200
TGATTTTCC	CTGAATCTGG	CCTCTATAC	TCACTTGGC	TTTTTTAAC	CGTTGGATAA	1260
TGATTTTCAT	AATAGCCTTT	CTAGTAAGAG	CTAGGACAAC	TAGCCGTTGG	TCCGTTGAC	1320
AGAGTAAACT	TCTGGCACAC	TCTTAATTTC	ATCGACAACC	GTGGTCAGTG	TAGAGAGGTT	1380
GGCAATACCG	AAGgACACAT	GGATATTAGC	AAACTTCATA	TCCTTGGTTG	GTTGGCATT	1440
GACCGTTGAA	ATATTCTTG	TTGTATTTGA	AAGAACCTGC	AGTACATCGT	TCAACAGTCC	1500
TGTACGGTTG	AGACCGTAGA	TATCGATATG	GGCCATATAC	TCCTTATTTG	AGCTAGGGTA	1560
CTGGTCTTCC	CATTCCACAT	CAAGGAGACG	TTGCTCGTAG	TTTTCTTGGG	CACGCAGGTT	1620
CATACAGTCC	ACACGGTGA	TAGCCACACC	ACGACCC TTG	GTAATGTAGC	CAACAATATC	1680
GTCACCAGGC	ACGGGGTTAC	AAACACTTAGC	AATCCGC ACT	AGGAGACCAG	AAGCACCTTC	1740
AATAACCACT	CCCCCCTCAT	GCTTGACCTT	GAGGGTTCT	TTATTTCAA	CCTTGACCTC	1800
GCCACCTTG	ACAAGCTCCT	CTGCCTCAGC	TTGGCCTTG	GCACGCTCTT	CCTCACGGCG	1860
TTCCCTTTCA	GTCAGACGGT	TAAAGACGGT	AATCGCACCG	ATTTCCCCAA	AACCAATGGC	1920
CGCAAAGAGG	GAGTCTTCTG	TCTTGTAACT	GGTCTTTGC	AGAACTTGAT	CCATGTGGCG	1980
CTTGTCCATA	AATTATTTTG	CCACATAGCC	ATTTCCTTG	AACTGAGCCA	TCAGCATCTC	2040
ACGACCC TTG	TTGACAGACA	ATTCC TTATC	TTGGTTTTA	AAGAACTGGC	GAATCTTATT	2100
GCGCGCCTTG	CTAGTCTTGA	CCATATTGAG	CCAGTCACGG	CTAGGTCAA	AGGAGTTCGG	2160
GTTGGCGATA	ATTTCAACCT	GATCCCCTGT	CTTTAACTTG	GTTGTCAGTG	GAACCATGCG	2220
GCCATTGACC	TTGGCACCAG	TTGCTTTTC	ACCGACCTTG	GTATGGATT	CGTAGGC AAA	2280
ATCAATCGGT	CCTGAATCTT	TGGGAAGGGA	ACGGACAGCT	CCATCTGGGG	TAAAAACGTA	2340
AATCTCCTCA	GCCAAATAGT	TTTCCTTAAC	AGAGTCCACA	AATTCCTTAG	CATCATCAGC	2400
CTGGTCTTGG	AGCTCCATCA	TCTCCTTGAT	CCAGTTCAT	CCAATAGCTG	ATTCC TTGCT	2460
GTAACTTGC	CCCTTATAC	CTTCTTATA	AGCCCAGTGA	GCCGCAACCC	CGTACTCAGC	2520
CACCTCGTGC	ATTTCC TTG	TTCGAATCTG	GAATTCAATC	GGCCCTTTG	GTCCATAAAC	2580
AGTCGTATGG	ATAGACTGAT	AACCATTGGC	CTTGC GGGTTG	GCGATATAGT	CTTTGAAGCG	2640

848

ACCTGGCATC GGTTTCCAAA ATTCAATGCAC GTAACCAAGC ATGGCATAAA CATCACTTTG	2700
GGTATCTAAA ATACAACGAA TAGCAATCAG ATCATAGATT TCCTCAAACC GTTTTCTCTT	2760
GTCCTGCATT TTGCGGAAAAA TTGAGTAAAT ATGCTTGGGA CGACCATAAA TCTTCCCTTT	2820
CAAGTGACGT TCTGTCGTAT ACTCCTCTAA TTTTGTGACT ACCTCATCCA CCAAGGCCTC	2880
ACGCTCCCTG CGCTTTCCCT TCATCATATG GGTAACTCTG TAAAACCTCG TTGGATTGAG	2940
ATAACGGAAA GACAAGTCTT CTAATTCCCA TTTGACACTG GAAATCCCCA AACGATGGGC	3000
AAGCGGGGCA TAGATTCCA TGGTTTCTTT GGAAATACGC TCCTGCTTGT CTNTTCAAG	3060
ATGTTTCAGG GTCCGCATAT TGTGCAAGCG GTCAGACAGT TTGACCAAA TAACGCGGAT	3120
GTCCTCAGAC ATGGCCATGA GCATCTTGCATG ATGATTTCC GCTAATTGCT CCTCGATCGA	3180
TTTGTACTCG ACCTTGCCAA GCTTGGTAAC TCCGTCAACA ATCATCCGCA CATCAGGACC	3240
AAACTCTCTT TCCAAATCGT CCAAAGTCGC ATCTGTATCT TCCACCACAT CATGCAAGAA	3300
TCCACAAGCT ACTGTTACAG CATCCAGCTT TAGCTTAGCT AAAATACCTG CCACCTGGAT	3360
AGGGTGAATG ATATAAGGCT CGCCTGATTT GCGATATTGA CCACTGTGGC ATTCAACAGC	3420
ATAGACCAAG GCCTTATGGA CAAATGAAC ATCCCTTCC GTTAAATATT CTNTGGTTAA	3480
AGCGACAAC TCTTCGCCCTG TTAAATTCAAC TTCTTTCGGC ATCTCTACTC TCCAATTCTT	3540
CCTACCATT TATCACTTTT TTAAGAATAT GAAAACCTAGA TTGGAACAGA ATAAGAAAAA	3600
AATAATTCAA AATTGCTTGA TAATTCTGAA TTATTGGTCC GTAATATACT ACGAAGTTAG	3660
ATTTTAAACT TAGGTGATAG AAGGAGAGAT AGAAGAACGG AAACCATATT GTAACCCAAA	3720
GACTTTCTGA CTTCCCCAAT TCCATTGAAG ATACGAAAGA TAAACGGTGG AACTCGTATC	3780
ACATACACTG GTACCTTGAC TGGATTTGG ATTAATACT AAATGAAAAT CAAAGAGCAA	3840
ACTAGGAAAC TAGCCGCAGG TTACTCAAAG CACCGCTTTG AGGTTGCAGA TAAAGTTGAC	3900
GCGGTTTGA GAGATTTTG AAGAGTATAA AAATCCTCAA GATACTTTCT TCTATCCTTT	3960
AGTTTATAAG GAGAATACCT ATGAAAAAAA CTGCTATTTC TATCTTGCT CTCCTAAATGT	4020
TAGGAGTTTG CTGCCTGTTCTATTCAGCC AGCAAAGCTA TAAAAAACAG TCGTTCAATA	4080
CTATGCTAAC GACCAGAAC TGCCCAGTAG GATAACTTAT AGTGAATATA GCGACAAATG	4140
AGAAGCCAAC TACGGTAGCA CTCTAACAT CACGTCTATC AAACAAGCTA ATGACGGAGT	4200
TTATGCAACC TATGAAGGGC AATTGACACC TTTCCAATAT TGATAAAATG ATAACCAGCC	4260
TGTCTTCATC TAGTCATGCT GGTTTTAAG TTCATTTAA ATCCTTACCT ATTCTCCCTA	4320
ACTGTGCTAT ACTTAATTAA TACTCAATGA AAATCAAAGA GCAAACCTAGA AAGCTAGCCG	4380
CAGGCTGTTCAAG CAGATAAAAGT TGACGGGTT TGAAGAGATT	4440

849

TTCGAAGAGT ATTAGTACAT TCTTTGAGAT TGGAGCTAGT ATGAAAATCC ATAAAACCGT	4500
GAATCCTGTT GCCTATGAAA ATACCTATTA TCTAGAAGGC GAAAAGCACC TCATCGTCGT	4560
CGATCCTGGT AGTCATTGGG AAGCCATTG TCAGACAATC GAGAAGATCA ACAAACCGAT	4620
CTGTGCTATT CTCTTGACCC ACGCCCATT TGACCATATC ATGAGTCTGG ACTTGGTTCG	4680
CGAGACGTTT GGCAATCCTC CTGTCTATAT CGCAGAGAGC GAAGCCAGCT GGCTCTACAC	4740
TCCTGTCGAT AATCTCTCCG GTCTCCTCG CCACGATGAT ATGGCAGATG TGGTCACAAA	4800
ACCTGCAGAA CACACCTTG TCTTTCACGA AGAATACCAA CTAGAGGAAT TTCGTTTAA	4860
GGTTCTACCG ACCCCAGGGC ACTCTATCGG TGGTGTTCC CTAGTCTTTC CTGATGCTCA	4920
TCTAGTCTTG ACGGGAGATG CTCTATTCCG CGAAACTATC GGACGGACCG ACCTTCCGAC	4980
TGGTAGCATG GAGCAACTCC TTCATAGTAT CCAGACCCAA CTCTTCACCC TACCAAACCA	5040
CGATGTCTAT CCAGGACATG GTCCAGCTAC TACTATCGCT CACGAAAAGG CCTTCAATCC	5100
CTTTTCTAG CAAGATGATG ACAATCGAAA TTAAAGTAAA CTATCCAGCA AATCTTCTA	5160
TTACAAAAGG CATCCTATCA AGGTTTCAC ACATGATTGG ATGCCTTTT TCTGATGACT	5220
AGATTTTTTG CATTACCAAA TAATCACGCG CTCCCTCTGGT GAACGCCACA TTCCGTCTCC	5280
TTCTTTGACA TCATAGTTG TAAAGAAATC GTCGAAGTTT GGTACTTGCA CATTGACACG	5340
GAGTTTGCT GGTGCGTGCA CATCGACGCT AGCCAAAAGT TTCATAAATT CTGGTCGACC	5400
TTTCATGCGC CAGATGCGAC CGAAGTTGTA GAAGAACTCT TCTGCTGAGA AGTCTGCTTC	5460
TCTCTTAGCT GCTTCAAGCG CTGCTGCGAT TCCTCCCAAG TCAGCCACGT TTTCTGATAC	5520
AGTCAATTAA CCGTTAATGG TTGCTCCATA AGAATCCTGT CCATCAAATT GGTCAATGAC	5580
TTTTTGTGTT TTCTCCTTGA AGGCAGCATA GTCGCTCTCT GTCCACCAAT CCTTGAGGCT	5640
ACCATTTTCG TCAAAGGAAG CCCCCTTAGT ATCAAAGGCG TGGGAAATT CATGGGCAAT	5700
CACTGCCCA ATACCACCGT AGTTAGCAGA AGATGACTGA TGCAAGTCAT AGAAAGGCAC	5760
CTGTAAAATG GCCGCTGGAA AGACAATCAG GTTCTCTGA GGATTGAGT AGGCATTGAC	5820
CATATGAGCA GGCATGCCCA ATTCCCTATA ATCTACAGGC TGGTTCCACT TACTCCAATC	5880
GTGCTTGATT TCCACACGCG CAAAGGCTAG AGCATTCTCA AAAAGACTGG CAGTTTCATT	5940
CACTACCTTA TCCTTGTAAC GTGCAGGCAA TTCTTCTGGT TAGCCAATAT AAGGTTTGAT	6000
CACATTGAGC TTCACCGATAG CCTGTTTACA GGTTTCTGGA GTGAGCCAGT CATTCTTAAG	6060
CAGACGCTCC TTATAAACAT CAATCATGGT TGCCACTTTT TTCTCCACAT CCGCCTTGGC	6120
TTCTGGAGAG AACTTCTCAC GGGCGTACCA AAGACCCAGG GCTTGCTTGA AAGGTTCTTG	6180

850

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TGCTAGATGA TAAGCTGCTT TGACCTTATC TTTTGCCCTCT GGAACTCCAG AAAGGGCACG       6240
GCTGTAGGCA CCAGACAAAA CACGGATATC CTCTGTTAAA TAGCTGGTTG AAAGATTGAC       6300
AACACTCAAA ATCAAGGTTG CTTAAGGAG AGACCAGGCT TCCTCACTGT AGAATTGCTC       6360
TGCTGCTTGC CAGAAACGTT CCTCGTCTAC AATAACCTTG TCTGGTAATT GCCCAATAAC       6420
TGCTTTGAAG AAGTCATCCA AAGGTAGGGC AGGCGCGAAT TTCTTGAAAT CTTCGTAAGA       6480
ATATGGATGA TAGAGTTAG CATATTCTGA ACTTTCTTCA TTAGAGAGCA CCACTGCCGC       6540
AACTCGGGCGG TCCAATTCAA GTCTTTTTC TAGCAAGTCT TCAATTCTT CATCAGAGAA       6600
ATCATAAGCC TTGAGGAGAT TTGCGCTGCT TTCTTCCAA AGAGTCAGA GCTCTTCGCG       6660
CTGAGGATGT TCTTCGCTAG AGTAGGTCGT ATCTGGCAAG ATTGTGCTTG GAGCGCTAGC       6720
CCATAGAACAA TTGATTCTAG CATCCATAAA GTCTGGCGAT ACACCAAAAG GAAGGAAGTT       6780
TGGTTTCCT GCAAGCTCAA ACTCTGCTAG TTTAGCTGTA AAATCCGCAA AAGTCTCCAA       6840
TTCTTGAAT TCTTAAGGA GTGGTAAGAC AGGTGTGATA CCGTCAGCTT CTCTCTTGTC       6900
AAAATCACGA ACTAGGCGGT GGTATTTGAC AAAGTTTCC AAGATAGCAT CCTCAGGCAC       6960
TTCTTCACCT GCTAACCACT TGTCTGTTGT CGCCAGCATE AGGTCTTCAA TTTCTGGTC       7020
TAAATCAACA AAACCTCCCTG TTGAGACTT ATCTGCTGGG ATTTCAAGCTG TCTGTTGCCA       7080
TTCTCCATTG ATAGCATCAT AAAAATCATC TTGATAACGT GTCATCTTGT TCTCGCTTTC       7140
ATTTGTATTT GCATTATCT TAACAAAAAT CG                                         7172

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(2) INFORMATION FOR SEQ ID NO: 121:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4518 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 121:

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CGGGAAGTTA TGCGATCTAG ACTTCGTTCC TGTACAGCTA CTTTCTCAGG TGGTCTTGTT       60
GTTTGTATGA GTTTGTTAG AGAGGATCTT TCTATGTCTT TCTTTCTTAT TTTTGTGTTA       120
TATGCTTTTC TGATTTCTTA TCTAATTAT GGTATTCTCA GACTAAAAAG GAAATACCGA       180
GTAGATGAAT AGCAAGGTTG TAGGTCTTCA GATTGATTGTT TAGCACTCTT GATAAAAGAG       240
TGCTAATTGTT TTGAGTTTT GTCTTGACAT TCTCTTCTAA GGGTGTATAA TAGAATCATG       300
AGTTAGCACT TGGATGCATT GAGTGCTAAT TGATCAGACA GAGAGGAGTG ATGAGATGGT       360
TACAGAGCGT CAGCAGGATA TTTAAATCT GATTATTGAC ATCTTTACCA AAACGCACGA       420

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851

ACCTGTCGGA	TCAAAAGCCT	TGCAAGAGTC	TATTAACTCT	AGCAGTGCAA	CCATTCTGAA	480
TGACATGGCG	GAACTAGAAA	AACAAGGGTT	GCTTGAGAAG	GCTCATACTT	CAA GTGGTCG	540
GATGCCAAGT	GTTGCTGGTT	TTCAGTACTA	TGTGAAACAC	TCACTGGATT	TTGACCCGGCT	600
GGCTGAAAAT	GAGGTATATG	AGATTGTCAA	AGCCTTTGAT	CAGGAATTCT	TCAAATTGGA	660
GGATATTCTG	CAAGAGGCTG	CTAACCTTACT	AACAGACCTG	AGTGGCTGTA	CGGTAGTGGC	720
ACTGGATGTT	GAGCCGAGCA	GGCACAGTTT	GACAGCCTTT	GATATCGTTG	TTTGGGGCA	780
ACATACAGCC	TTGGCGGTAT	TTACCCCTAGA	CGAGTCGCGA	ACGGTTACTA	GTCAGTTCT	840
GATTCCAAGG	AACTTCTTGC	AGGAGGATTT	GCTGAAACTG	AAGAGCATCA	TTCAGGAACG	900
TTTCCTCGGT	CACACCGTTT	TAGATATTCA	CTACAAGATT	CGGACGGAGA	TTCCGCAGAT	960
TATCCAGCGT	TACTTTACAA	CAACGGATAA	TGTCATCGAT	CTCTTGAAC	ACATCTTAA	1020
GGAAATGTTT	AACGAAAACA	TTGTGATGGC	GGGCAAGTC	CATCTCTGA	ATTTTGCCTA	1080
TCTAGCAGCC	TATCAGTTCT	TTGACCAACC	GCAAAAGGTG	GCCTTGGAGA	TTCGTGAGGG	1140
GTTGCGTGAG	GATCAGATGC	AAAATGTTCG	TGTTGCAGAC	GGTCAAGAGT	CCTGTTAGC	1200
TGACCTAGCG	GTAATCAGTA	GTAAGTTCCCT	CATTCCCTAT	CGGGGAGTTG	GAATTCTAGC	1260
CATTATCGGT	CCAGTTAACATC	TGGATTACCA	ACAGCTAAC	AATCAAGTCA	ATGTGGTCAA	1320
CCGTGTTTG	ACCATGAAGT	TGACAGATTT	TTACCGCTAC	CTCAGCAGTA	ATCATTACGA	1380
AGTACATTA	GATTGAAATC	ATTAAAGGAG	GCGAACATGG	CCCAAGATAT	AAAAAATGAA	1440
GAAGTAGAAG	AAGTTCAAGA	AGAGGAAGTT	GTGAAAACAG	CTGAAGAAC	AACTCCGTAA	1500
AAAGTCTGAGT	TGGACTTGGC	AAATGAACGT	GCAGATGAGT	TCGAAAACAA	ATATCTTCGC	1560
GCTCATGCAG	AAATGAAAAA	TATCCAACGC	CGTGCCAATG	AAGAACGTCA	AAACTTGCAA	1620
CGTTATCGTA	GCCAGGACTT	GGCAAAAGCA	ATCTTACCAT	CTCTTGACAA	CCTTGAGCGT	1680
GCACATTGCA	TTGAAGGTTT	GACAGATGAT	GTGAAGAAGG	GCTTGGGGAT	GGTGCAAGAA	1740
AGCTTGATTC	ACGCTTGAA	AGAAGAAGGA	ATTGAAGAAA	TCGCAGCAGA	TGGCGAATTT	1800
GACCATAACT	ACCATATGGC	CATCCAAACT	CTCCCAGCAG	ACGATGAACA	CCCAGTAGAT	1860
ACCATCGCTC	AAGTCTTC	AAAAGGCTAC	AAACTCCATG	ACCGCATCCT	ACGCCAGCA	1920
ATGGTAGTGG	TGTATAACTA	AGATATAAAAG	CCCGTAAAAAA	GCTCGCAGTA	AAAATAGGAG	1980
ATTGACGAAG	TGTTCGATGA	ACACAAGAAA	ATCTATCTTT	TTTACTCAGA	GCTTAGGGCG	2040
TGTTCGATTC	GGCAATTCTG	ACGGTAGCTA	AAGCAACTCG	TCAGAAAACG	GCAATCGCTA	2100
TGGCGTTGC	CTAGCTTCCT	TACTAACTCG	TCGTCGAAAT	AAAATCGATT	TCGACTCCCTC	2160

852	
GTGTCGCAAT TTACATAATA GAAAACTTGT CCGAAACGAC AATAAACTAT GAAGAAAGAT	2220
AAAATATGTT TGGCTTGTA ATAGTGAGCG AAGCGAACCA AACACGATAC TCTTCGCCGT	2280
GGCGCTATTG GCGCAAATTT TGAGACCTTA GGCTCAAAGT TTAGTCAAAG AGATTGACGA	2340
AGTCAAGCTC TGACGGCGTC GCCACTGTCG CCACCTTAAGA AGAGTATCAA AAAGAAAAAT	2400
AGAAAATTAA CTAACAAGGA GAAAAACACA TGTCTAAAAT TATCGGTATT GACTTAGGTA	2460
CAACAAACTC AGCAGTTGCA GTTCTTGAAAG GAACTGAAAG CAAAATCATC GCAAACCCAG	2520
AAGGAAACCG CACAACCTCA TCTGTAGTCT CATTCAAAAA CGGAGAAATC ATCGTTGGTG	2580
ATGCTGAAA ACGTCAAGCA GTTACAAACC CAGATACAGT TATCTCTATC AAATCTAAGA	2640
TGGGAACCTTC TGAAAAAGTT TCTGCAAATG GAAAAGAATA CACTCCACAA GAAATCTCAG	2700
CTATGATCCT TCAATACCTG AAAGGCTACG CTGAAGACTA CCTTGGTGAG AAAGTAACCA	2760
AAGCTGTTAT CACAGTTCCG GCTTACTTCA ACGACGCTCA ACGTCAAGCA ACAAAGACG	2820
CTGGTAAAAT TGCTGGTCTT GAAGTGAAC GTATTGTTAA CGAACCAACT GCAGCAGCTC	2880
TTGCTTATGG TTTGGACAAG ACTGACAAAG AAGAAAAAAAT CTTGGTATTT GACCTTGGTG	2940
GTGGTACATT CGACGTCCT ATCCTTGAAAT TGGGTGACGG TGTCTTCGAC GTATTGTCAA	3000
CTGCAGGGGA CAACAAACCTT GGTGGTGACG ACTTTGACCA AAAAATCATT GACCACTTGG	3060
TAGCAGAATT CAAGAAAGAA AACGGTATCG ACTTGTCCTAC TGACAAGATG GCAATGCAAC	3120
GTTTGAAAGA TGCGGCTGAA AAAGCGAAGA AAGACCTTTC TGGTGTAAC TCAACACAAA	3180
TCAGCTTGCC ATTTATCACT GCAGGTGAGG CTGGACCTCT TCACTTGAA ATGACTTTGA	3240
CTCGTGCAGAA ATTTGACGAT TTGACTCGTG ACCTTGTGAA ACACGTTACAAA GTTCCAGTT	3300
GTCAAGCCCT TTCAGATGCA GGTGGAGCT TGTCAAGAAAT CGACGAAGTT ATCCTTGTG	3360
GTGGTTCAAC TCGTATCCCT GCCGTTGTTG AAGCTGTTAA AGCTGAAACT GGTAAAGAAC	3420
CAAACAAACCT AGTAAACCCCT GATGAAGTAG TTGCTATGGG TGCGGCTATC CAAGGTGGTG	3480
TGATTACTGG TGATGTCAAG GACGTTGTCC TTCTTGATGT AACGCCATTG TCACTTGGTA	3540
TCGAAACAAAT GGGTGGAGTA TTTACAAAAC TTATCGATCG CAACACTACA ATCCAAACAT	3600
CTAAATCACA AGTCTTCTCA ACAGCAGCG ACAACCAACC AGCCGTTGAT ATCCACGTT	3660
TTCAAGGTGA ACGCCAATG GCAGCAGATA ACAAGACTCT TGGACGCTTC CAATTGACTG	3720
ATATCCCAGC TGCACCTCGT GGAATTCCCTC AAATCGAAGT AACATTTGAC ATCGACAAGA	3780
ACGGTATCGT GTCTGTTAAG GCCAAAGACC TTGGAACCTCA AAAAGAACAA ACTATTGTCA	3840
TCCAATCGAA CTCAGGTTG ACTGACGAAG AAATCGACCG CATGATGAAA GATGCAGAAC	3900
CAAACGCTGA AGCCGATAAG AAACGTAAG AAGAAGTAGA CCTTCGTAAT GAAGTAGACC	3960

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AAGCAATCTT	TGCGACTGAA	AAGACAATCA	AGGAAACTGA	AGGTAAAGGC	TTCGACGCAG	4020
AACGTGACGC	TGCCCAAGCT	GCCCTTGATG	ACCTTAAGAA	AGCTCAAGAA	GACAACAAC	4080
TGGACGACAT	GAAAACAAAA	CTTGAGCAT	TGAACGAAAA	AGCTCAAGGA	CTTGCTGTTA	4140
AACTCTACGA	ACAAGCCGCA	GCAGCGAAC	AAGCTCAAGA	AGGAGCAGAA	GGCGCACAAAG	4200
CAACAGGGAA	CGCAGGGCAT	GACGTCGTAG	ACGGAGAGTT	TACGGAAAAG	TAAGATGAGT	4260
GTATTGGATG	AAGAGTATCT	AAAAAAATACA	CGAAAAGTTT	ATAATGATTT	TTGTAATCAA	4320
GCTGATAACT	ATAGAACATC	AAAAGATTTT	ATTGATAATA	TTCCAATAGA	ATATTTAGCT	4380
AGATATAGAG	AATTATATTA	GCTGAACATG	ATAGTTGTAT	CAAAAATGAT	GAAGCGGTAA	4440
GGAATTTTGT	TACCTCAGTA	TTGTTGTCTG	CATTTGTATC	GGCGATGGTA	CCGTATCTGA	4500
CGAACGTTCA	GCTTATAT					4518

(2) INFORMATION FOR SEQ ID NO: 122:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8145 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 122:

TGCTATTTTC	GATTCCCTTG	GGCGTTTGA	TTGCCTTGC	CTTGCAAGTC	CATTGGAAGC	60
CCCTCCATTA	TCTGATTAAC	ATTTACATCT	GGGTTATGCG	AGGAACCCCC	TTACTCTTGC	120
AACTGATTTT	TATCTATTAT	GTGCTCCCAA	GTATTGGAT	TCGTTAGAC	CGCCTTCCTG	180
CAGCTATTAT	TGCCTTGTT	CTCAACTATG	CAGCTTACTT	TGCAGAAATT	TTCCGTGGGG	240
GAATTGACAC	TATTCCAAGA	GGACAGTATG	AGGCCGCCAA	GGTCTTGAAG	TTTAGCCCTT	300
TTGACAGAGT	GCGCTATATT	ATCTGCCCA	AAGTGCACAA	GATCGTTCTT	CCTAGTGTCT	360
TTAATGAAGT	TATGAGTTG	GTCAAGGATA	CTTCTTGGT	CTATGCTCTC	GGAATTCAG	420
ACCTTATCTT	GGCTAGTCGA	ACAGCTGCTA	ACCGCGATGC	TAGTCTAGTT	CCTATGTTCT	480
TGGCAGGAGC	CATTTATTTG	ATTTGATTG	GGATTGTGAC	AATTATTTCC	AAAAAAAGTTG	540
AGAAGAAAGTA	TAGTTATTAT	AGATAGGAGG	CTGCCATGTT	AGAATTACGA	AATATCAATA	600
AAGTCTTGG	AGACAAACAA	ATCCTGTCTA	ATTCAGTCT	AAGTATTCC	AAAAAGCAAA	660
TCCTGGCTAT	CGTTGGACCT	TCTGGTGGAG	GTAAGACAAC	TCTTTACGT	ATGCTTGCAG	720
GTCTTGAAAC	CATTGATTCA	GGGCAAATCT	TTTATAATGG	ACAACCTTA	GAGCTGGATG	780

854	
AATTGCAGAA GCGCAATCTA CTGGGATTTG TCTTCCAAGA TTTTCAACTA TTTCCATC	840
TATCAGTTCT GGAAAATTG ACTTTATCGC CTGTGAAGAC CATGGGAATG AAGCAGGAAG	900
AGGCTGAGAA GAAGGCGAGT GGACTCTTGG AACAGTTAGG ACTAGGAGGA CACGCAGAGG	960
CCTATCCTT CTCACTATCT GGTGGCAAAGCAGCGGGT GGCTTGCG CGTGTATGA	1020
TGATTGACCC AGAAATCATT GGCTACGATG AACCAACTTC TGCCCTGGAT CCAGAATTAC	1080
GTTTGGAACTT GGAGAACGTA ATCTTGCAAA ATAGGGAACT TGGGATGACC CAGATTGTGG	1140
TTACCCATGA TTTGCAGTTT GCTGAAAATA TCGCAGATGT ATTATTGAAA GTAGAACCTA	1200
AATAGGAGGA AAAATGGATG AAAAAATGGA TGCTTGTATT AGTCAGTCTG ATGACTGCTT	1260
TGTTCTTAGT AGCTTGCGG AAAAATTCTA GCGAAACTAG TGGAGATAAT TGGTCAAAGT	1320
ACCAGTCTAA CAAGTCTATT ACTATTGGAT TTGATAGTAC TTTGTTCCA ATGGGATTG	1380
CTCAGAAAGA TGGTTCTTAT GCAGGATTG ATATTGATT AGCTACAGCT GTTTTGAAA	1440
AATACGGAAT CACGGTAAAT TGGCAACCGA TTGATTGGGA TTTGAAAGAA GCTGAATTGA	1500
CAAAAGGAAC GATTGATCTG ATTTGGAATG GCTATTCCGC TACAGACGAA CGCCGTGAAA	1560
AGGTGGCTTT CACTAACTCA TATATGAAGA ATGAGCAGGT ATTGGTTACG AAGAACATCAT	1620
CTGGTATCAC GACTGCAAAG GATATGACTG GAAAGACATT AGGAGCTCAA GCTGGTTCAT	1680
CTGGTTATGC GGACTTTGAA GCAAATCCAG AAATTTGAA GAATATTGTC GCTAATAAGG	1740
AAGCGAATCA ATACCAAACC TTTAATGAAG CCTTGATTGA TTTGAAAAAC GATCGAATTG	1800
ATGGTCTATT GATTGACCGT GTCTATGCAA ACTATTATTT AGAAGCAGAA GGTGTTTAA	1860
ACGATTATAA TGTCTTACA GTTGGACTAG AAACAGAAGC TTTTGCCTT GGACCCGTA	1920
AGGAAGATAAC AAACCTGGTT AAGAAGATAA ATGAAGCTTT TTCTAGTCTT TACAAGGACG	1980
GCAAGTTCCA AGAAATCAGC CAAAAATGGT TTGGAGAAGA TGTAGCAACC AAAGAAGTAA	2040
AAGAAGGACAA GTAAGATAAA ATAGTGGCTG AAAC TGCGTT TTGATTAGCA AAACGTAGTT	2100
TTTTTGTAATCTAGGAAAA CGATAATAGC GATTGAATAT GGATAATTGA ATATGGAATA	2160
GCCCCACTGTG ATTTCTAAAA CATTGTAAA AATTGATTG ACTTCCAAAA TTAAATGTT	2220
CTGTAATGAA ATACTGATGT AACTGTTTA GGAACAATAA AACGCATAAT ATCAAGGTTT	2280
TTGCACCTTA CATTATGCGT TTTTGTGATT TTAAGACTTG TTAGCTGATT TTTTACAATC	2340
CTGCGAAATC TTTGATTTCT TGTGCTGACA TTGAAGAGTC GCAACGGACG TTGATTGTC	2400
CATCTGTAAT ATGAACAAAA CCTGGTACAG TTGGGATTCC ATAGCGTGAG CGGAATGCTT	2460
GCAAATCATT GAGTTGGCTT GGTTCTTCAC TATTGATGAA GTAAATGTGA GCTTTGGTTT	2520
CAGCTACGAC ACCTGACAAT GTACCTGCAA ATTTACGGCA GTAAGGGCAA GTTTTGCGAC	2580

855

CGATAAAGAA GGTTGCAGTT TCTTTTTAT CAAGAGCTTC TTGCGCACGC ACAACTGTAG	2640
TGACTTCAAG GTCTTGATG TTATCTAAAA ATTGTTCCAT GAGATTACCT CGCTTTCATT	2700
GATAAGTCTA GTATGCCATA AAGTTCTAA AATTGCTTAG ATTTGATACG AAAAAAGATG	2760
AGGTTGGTTG GTCTCATCTT TTATAGGTCT TTATTTACA AATGCATTGA TTTCTGCTTC	2820
GATGTTAGCA ATCTTAGCTT GTGATTCTTC GTTGGTTCC CCTACAACGT CAATGTAGAA	2880
CTTGATTTT GGTTCTGTAC CTGAAGGGCG AACGGCAATC CATGAACCGT CAGCAAGTGT	2940
GTATTTCAAC ACATCACTTG GAGGAGTTGT CAAGTTGTAA ACAGTACCGT CAGCAACAGT	3000
AGCAGTTTGT GCCTTGAAGT CTTCTACGAC AGTGTAGCT GTTGCCTTCC ATTCTGTTGG	3060
AGCATTGTTG CGGAATTAG CCATAATCGC TTTGATTTGT TCAGCACCCT GAACACCTGA	3120
AAGAGTAACA GAGATTGTTT TTTCTGCGTA GTAGCCATAT TCTTTATAGA TTTCTTCGAT	3180
ACCGTCAGCA AGTGTCAAAC CACCGAGAACG GTAGTAGGCA GCAAGTTCAAG CAACTACAAG	3240
AACGGCTTGG ATGGCATCTT TATCACGTAC AAATGGTTA ATCAAGTAAC CGAACGTTTC	3300
TTCAAATCCC ATCATGTAAG TGTGGTTGTG TTTTCTTCG AATTCTGGAA TTTTTTCAGC	3360
GATAAAATTG AAACCTGTCA AGACGTTGAA CATAGTTGCG CCGTAGCTTT CAGCAATCTT	3420
CGTTACCAAG TCAGTTGAAA CGATAGATTT GCAGAGAGCG GCATTTTCAG GAAGAGTTCC	3480
AGCGTTTTG TGAGCTTCCA AGATGTATTT AGCCATGATA GCACCGATTG GTTACCTGA	3540
AAGGTTGAGG TAGCTACCCT CTTTTGAAAG AACTCAACA CCAACACGGT CAGCGTCTGG	3600
GTCAGTTGCG ACAAGAACAT CTGCACCAAC TTGACGACCA AGTTCTTCAG CAAGGGCAAA	3660
GGCTGCTTGG CTTTCTGGGT TTGGAGATGT TACAGTTGAA AAGTCTGGGT CAGCAGTTGC	3720
TTGCGCTTCA ACAACTTGAA CAGAGTCAAA TCCTGCTTGG GCAAGAGCAC GACGAGCCAA	3780
CATTTCACCA GTACCATGAA GTGGTGTGTA GACAATCTTC ATGTCTTAC CAAATTCTTC	3840
AATCAAGGCT GGGTTGATGT TTATGTCCTT AACCTTTA AGGTATTCTA TGTCAACAGC	3900
TTCGCCGATA ACTTCAATCA AGCCAGAACG TTTTCAGTT TCCACATCAG CAACTCAAC	3960
TGCAAATGGG TTTTCGATTG CACGGATATA AGTAGTCAAA GCGTCCGCAT CGTGTGGAGG	4020
CATTTGTCCA CCGTCTTCAC CGTAAACCTT GTAACCGTTA AATGGAGCAG GGTTGTGGCT	4080
GGCTGTGACC ATGATACCTG CGAAACAGTT GAGATGACGA ACTGCAAATG ATAGTTCTGG	4140
AGTCGGACCA AGGCTTCAA ATACGTAAGA TTTGATGCCG TGTTTAGCAA GAACTGCCGC	4200
AGATTCAAAG GCAAACTCAG GTGAGAAGTG ACGGCTATCG TAGGCAATTG CTACACCGCG	4260
TTCTTTCTCG TTTCCACCTT TTGACTCAAT CAAACGAGCC AATCCTTCAG TAGCTTGGCG	4320

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AACAAACGTAG ATGTTGATAC GGTTTGTACC AGCACCAACC AAGCCACGCA TACCTGCAGT	4380
ACCAAATTCA AGATTTGTAT AGAAGGCATC TTCCCTTAGTT TTTTCGTCCA TATTTTCCAA	4440
ATCTTGACGA AGGTAGTCAC GAAGCTCCAC AAAATCAACC CATTCTGGT AATTTCTTG	4500
GTAAGACATT CAAATTCTCC TTTATTTTA AACACATTAA TCAGTTAAC TATATCATTT	4560
TTTTTAGTTT TAGTAAAACC TTATCTGCTT CGAACATCTC TTCAAACCGAG GTCAGATTGA	4620
ATTTTGGGGT TATATGATGT TGAGGCTAGG AAAAATTCAA TTTCAGTAAA AAAAGTAAGT	4680
CTTCTCATAA CAAAACATTG ATATAGTTAC TTAGTTAAC ACAAGCATAT TATAATAAAG	4740
CTATGGCATA TAGTACTGAT TTTAACACGC GAGCATTAGA TTACATCAA GAGGGGCACA	4800
GCCATGTCGA GCCAGCCAAG TTTTTTGTG TTGGCGTCAG AACTCTCTC ACGTGGAAA	4860
AGAAAGACGT GAACAAGAAC ACATAGAGAG GAAAAAGCGA GTCGTAAAA ACCGAAAGAT	4920
TCCTTTAGAG GAATTGAAAG CCTTTGTAGA GGCTCATCCA GATGCTTTT TACGGGAAAT	4980
TGCGGCACAT TTTGATTGTG CTGTTCCCTC AGTATGGCA GCTTTAAAGC AGATTAAGGT	5040
CACTTTAAAA AAAGATGACG AGCTTTAACCG AACAAAGACCC AGAAAAGTAG CCTTATTCT	5100
TAAGAATTAA AATAGTTAA AGCACCTAGC ACCTGTTTAT ATTGATGAAA CAGGAATCGA	5160
CCGCTATCTC TATCGTCCTT ATGCAGGGC TCCTAGAGG GAGAAAGTCT ATGAAAAGAT	5220
TAGCGGACGT CGTTTGAGC GAACTCAAT TGTTGCAGGA CAAGTAGACG GAGAGTTAT	5280
AGCTCCCAGT ATTTACAAGA AAACCATGAC AAGCGATTTT TTTGTGGAGT GGTCAAAAC	5340
GCAACTCCTA CCTGCTTGA AGACACCTCA TGTTATTGTC ATGGCAATG CTGGTTTCA	5400
TCCCAAGAAC ATTTGGATG AACTCTGCAT CCAAGATAAA CACTTTCT TACCTCTACC	5460
ACCTTATTCA CCGGATTGTA ATCCTATTGA GCAAGCTGG GCTATCTGA AAAAGAAAGT	5520
GACGGATGTA TTAAGGGAAG TTCCAACAT TTTGAATGT TTGGAATGCT TTTTAAAC	5580
TAGATGACTA TAACGGTTCT AAAGGAACCT ATCGAGTAGT CATTAAACT AAGGATACTG	5640
CTGGTTAAGA GAAGACGGTA TACAATCAAA CCATTACCG TGAGGCCAA ATCGTTCAGA	5700
ATGAAGACTT GTATCAGAAT GAAGACTTGT ATAAGAAAGG TTTGAATGTT GAACTTGC	5760
ACCAACAAAT TAAGGGATT TTTGAAGCAG AGTTAAAAAA TCGTATTAAT GGAGTTCTTA	5820
ATACTAAAAT AAAAATAGT ACATTAATC GTGAAATAA AAAAATATA CACCAGAGCA	5880
ACAAAAACTC CATGATCAAT TTGAAGCAGA AGCAACGGAA GATGCTAAA AACAGGC	5940
TATTGTGTTG AATGTTGACC AGGATTCAT GAGCATATCT AAGTCTAATA AAAGTGGTTC	6000
AGACTGGAAG AAAACTTCA CAGTGAGGAT AACCAATAGG CTAGCAAATG ACTTGAATAA	6060
TGTCTTGAAA CAGGTTGATA AAGATACTCC TAATACCCCA ACTTGGCTAA ACTCAGCTGC	6120

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TTCTAAAGCT	AAAGATGATG	ACAGAGTATA	TAAAATCTG	AAGACTCTTA	TACCAGGAGA	6180
AAATTACCTA	TCATGTTAACG	GATAATCAGC	TAGAAGTAGA	AACAGATAAA	TACACATATA	6240
CTGCCGCTAG	AAATGGTAGT	AAGGAAGTTG	GTATTCAAGA	GTCAGATATA	GCAGCAACTC	6300
TAAGTGCCGA	TGAATATAAT	TCTAATCGCC	AAACTTTGA	GAGAGAATAC	AAATACAAAA	6360
GCAAATGCC	TTAATAATGG	TTGGGCTAGA	TCTGGTTCTG	AAGAGTTCAA	AAAGTTCTCC	6420
CACTTTGTAG	GGGTAGACAA	AGGGATTGTG	CGAACGAATG	TACTGACTGG	TAAAAAACTA	6480
TCTGATAAGA	TTAGGAAAGA	AGTGGGCTCT	GGAGATAGCA	AACTAGGAAA	AGGCGGCTAT	6540
TTCTCTACTG	GGGATGTTCT	ATTAGGAAA	GATGTTGTTT	CTTATACCGT	ACAAGTATTT	6600
TCAGAGAATA	ATGAAAGAGT	AGGAGTAAAC	ACTCAAAGTC	ACCGTGTTCA	GTATAATCTC	6660
CCAATTCTAG	CTGACTTTTC	AGTCATCCAA	GATACTGTGG	AACCACACG	AACC GTTGT	6720
GAAAAAAATCA	TTCCAAAACT	AAATATTCCC	GAAGAAGAGA	AAGGGAAAAT	AACCGAAGAA	6780
ATCAAGAAAA	AGAAAAAAAC	CTCAGAATTG	GCAGAACTAA	TCTCAGAAAA	TGTGAAAGTT	6840
CGCTATGTTG	ATGAACAAGG	GC GTTGCTA	TCATTGAAA	ATGATACTGG	AATTGGAGAA	6900
AAAGAAAGTG	ACGGAACCTA	CATTACCAAT	AAAAAACAAAC	TGATTGGTAC	CAGCTATAAT	6960
GTCACAGATA	AAAAACTCG	TAGCATGACT	ACTACTGACG	GAAAATATTA	TACTTTAAA	7020
GAAGCAGATA	CAAATTCTGC	AAAGTTAACT	GGGAATATTG	TAAGCGAAGG	TAGAACAGTG	7080
ACCTTAGTTT	ATAGAGAAAG	CGAAGCGCCA	ACCACTGCTA	CAGTAACAGC	CAATTACTAT	7140
AAAGAAGGTA	GGCAAGAGAA	GTTGGTAGAG	TCTGTTATAA	AAGCTGATTT	AGCGATAGGT	7200
TCTGAGTATA	CCACAGAATC	AAAAACTATT	GAAGGGAAAA	CAACAACTGA	GGACAAAGAA	7260
GACCGAGTTA	TCACAAGGAA	AACAACATAC	ACCTTGGTAG	CAACTCCTGA	AAATGCGTAC	7320
CAGAAGACGG	TGCAACAGTT	GACTATTACT	ACCGTGAGAA	TGTTGAGGAA	ACAGTGGTTC	7380
CCAAAACAGC	AACCTCTACT	GAGACGAAGA	CTATAACGCG	TATCATTAT	TACGTTGATA	7440
AAGTTACGAA	CCAAAATGTA	AAAGAAGATG	TTGTTCAACC	TGTAACCTTA	AGCCGTACAA	7500
AAACTGAGAA	CAAGGTCAACG	GGAGTTGTAAC	CCTACGGTGA	ATGGACAACA	GGAAACTGGG	7560
ACGAGGTTAT	ATCTGGTAAG	ATTGACAAGT	ACAAAGATCC	AGATATTCCA	ACAGTTGAAT	7620
CACAAGAAGT	TACGTCAACG	TCTAGTGATA	AAGAAATAAC	GGTAAGGTAT	GACCGTTTAT	7680
CAACACCAGA	AAAACCAATC	CCACAACCAA	ATCCAGAGCA	TCCAAGTGT	CCGACACCAA	7740
ACCCAGAACT	ACCAAATCAA	GAGACTCCAA	CACCAGATAA	ACCAACTCCA	GAACCAGGTA	7800
CTCCAAAAAC	TGAAACTCCA	GTGAATCCAG	ACCCAGAACT	TCCGACTTAT	GAGACAGGTA	7860

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AGAGAGAGGA ATTGCCAAC ACAGGTACAG AAGCTAATGC TACCTGGCT AGTGCTGGTA	7920
TCATGACCTT GTTAGCTGGT CTAGGATTAG GATTTTCAA GAAAAAAAGAA GATGAAAAT	7980
AATAGATTT AGAATCTAGG ACCAGGAAA AGCTCACAGA TGTGGGCTTT TTTCTGGTT	8040
TTGAGAACCGA GGTCTTCGT AAAGAATAAA AACGCTTACA AGTCTGTTGA ACTGGGAAAC	8100
TATGAATCCT ATTTTTTAA AAATATTCC AGAAATCACT TGCGG	8145

(2) INFORMATION FOR SEQ ID NO: 123:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8697 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 123:

CGGTACCGGG AACGATACTT AGTCTAATTG TGCACCTTT CCATGTATGG TAAAGGTTTT	60
TCTTTTTTA AAAAGGAAA CGAGAACAGG AGGTTCTTAT GAAAGCAAGC ATTGCCTTGC	120
AAGTTTTACC CCTAGTACAG GGGATTGATC GGATAGCTGT TATTGATCAG GTCATTGCTT	180
ATCTGCAwAC TCAAGAACGT ACGATGGTAG TGACACCATT TGAAACGGTC TTGGAAGGGG	240
AGTTTGATGA GCTTATGCGC ATTCTAAAAG AACGCGCTGGA AGTGGCAGGG CAGGAGGCAG	300
ACAATGTCTT TGCCAATGTC AAAATAATG TAGGAGAGAT TTTAAGTATT GATGAGAAC	360
TTGAGAACGTA TACTGAGACG ACACATTAGT CTATTGGCT TTCTCGGAGT ATTGTCAATC	420
TGGCAATTAG CAGGTTTCT TAAACTTCTC CCCAAGTTA TCCTGCCGAC ACCTCTTGAA	480
ATTCTCCAGC CCTTTGTCG TGACAGAGAA TTTCTCTGGC ACCATAGCTG GGCGACCTTG	540
AGAGTGGCTT TACTGGGCT GATTTGGGA GTTTGATTG CCTGTCTTAT GGCTGTGCTC	600
ATGGATAGTT TGACTTGGCT CAATGACCTG ATTTACCTA TGATGGTGGT CATTCAAGACC	660
ATTCCGACCA TTGCCATAGC TCCTATCCTG GTCTTGTGGC TAGGTTATGG GATTTGCCC	720
AAGATTGTCT TGATTATCTT AACGACAACC TTCCCCATCA TCGTTAGTAT TTTGGACGGT	780
TTTAGGCATT GCGACAAGGA TATGCTGACC TTGTTTAGTC TGATGCGGGC CAAGCCTTGG	840
CAAATCCTGT GGCATTTAA AATCCCAGTT AGCCTGCCTT ACTTTTATGC AGGTCTGAGG	900
GTCAGTGTCT CCTACGCCTT TATCACAACG GTGGTATCTG AGTGGTTGGG AGGTTTGAA	960
GGTCTTGGTG TTTATATGAT TCAGTCTAAA AACTGTTTC AGTATGATAC CATGTTGCC	1020
ATTATTATTC TGGTGTGCGAT TATCAGTCTT TTGGGTATGA AGCTGGTCGA TATCAGTGAA	1080
AAATATGTGA TTAAATGGAA ACGTCGTAG AATTAGAATG TTTCTGAAAA AGAAAAGAGG	1140

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AAATCAAAAT	GAAGAAAACA	TGGAAAGTGT	TTTTAACGCT	TGTAACAGCT	CTTGTAGCTG	1200
TTGTGCTTGT	GGCCTGTGGT	CAAGGAACTG	CTTCTAAAGA	CAACAAAGAG	GCAGAACTTA	1260
AGAAGGTTGA	CTTTATCCTA	GACTGGACAC	CAAATACCAA	CCACACAGGG	CTTTATGTTG	1320
CCAAGGAAAA	AGGTTATTTC	AAAGAAGCTG	GAGTGGATGT	TGATTTGAAA	TTGCCACCAG	1380
AAGAAAGTTC	TTCTGACTTG	GTTATCAACG	GAAAGGCACC	ATTTGCAGTG	TATTTCCAAG	1440
ACTACATGGC	TAAGAAATTG	AAAAAGGAG	CAGGAATCAC	TGCCGTTGCA	GCTATTGTTG	1500
AACACAATAAC	ATCAGGAATC	ATCTCTCGTA	AATCTGATAA	TGTAAGCAGT	CCAAAAGACT	1560
TGGTTGGTAA	GAAATATGGG	ACATGGAATG	ACCCAACTGA	ACTTGCTATG	TTGAAAACCT	1620
TGGTAGAATC	TCAAGGTGGA	GACTTTGAGA	AGGTTGAAAA	AGTACCAAAT	AACGACTCAA	1680
ACTCAATCAC	ACCGATTGCC	AATGGCGTCT	TTGATACTGC	TTGGATTTAC	TACGGTTGGG	1740
ATGGTATCCT	TGCTAAATCT	CAAGGTGTA	ATGCTAACTT	CATGTA	TTGAACTATG	1800
TCAAGGAGTT	TGACTACTAT	TCACCA	CGTAA	CAACGACTAT	CTGAAAGATA	1860
ACAAAGAAGA	AGCTCGAAA	GTCATCCAAG	CCATCAAAA	AGGCTACCAA	TATGCCATGG	1920
AACATCCAGA	AGAAGCTGCA	GATATTCTCA	TCAAGAATGC	ACCTGA	ACTC AACGAAAAAC	1980
GTGACTTTGT	CATCGAATCT	CAAAAATACT	TGTCAAAGA	ATACGCA	AGC GACAAGGAAA	2040
AATGGGGTCA	ATTGACGCA	GCTCGCTGGA	ATGCTTTCTA	CAAATGGGAT	AAAGAAAATG	2100
GTATCCTTAA	AGAAGACTTG	ACAGACAAAG	GCTTCACCAA	CGAATTG	TG AAATAATGAC	2160
AGAAATTAGA	CTAGAGCACG	TCAGTTATGC	CTATGGTCAG	GAGAGGATT	TAGAGGATAT	2220
CAACCTACAG	GTGACTTCAG	GCGAAGTGGT	TTCCATCCTA	GGCCCA	AGTG GTGTTGGAAA	2280
GACCACCCCTC	TTTAATCTAA	TCGCTGGGAT	TTAGAGGTT	CAGTCAGGGA	GAATTGTCCT	2340
TGATGGTGA	GAAAATCCC	AGGGCGCGT	GAGTTATATG	TTGCAA	AGG AGG ATCTGCTCTT	2400
GGAGCACAAG	ACGGTGCTTG	GAAATATCAT	TCTGCCCTC	TTGATTCA	AA AGGTGGATAA	2460
GGCAGAAGCT	ATTCCCCGAG	CGGATAAAAT	TCTTGC	GACCC	GACC TGAAG	2520
AGACAAGTAT	CCTCATGAAC	TTAGCGGTGG	GATGCGCCAG	CGTGTAGCCT	TACTCCGGAC	2580
CTACCTTTT	GGGCACAAGC	TCTTCTCTT	AGATGAGGCC	TTAGCGCCT	TGGATGAGAT	2640
GACAAAGATG	GAAC	CTTGGTATCT	TGAGATTCA	AAGCAGTTGC	AGCTAACAAAC	2700
CCTGATCATC	ACGCATAGTA	TTGAGGAGGC	CCTCAATCTC	AGCGACCGTA	TCTATATCTT	2760
GAAAATCGC	CCTGGGCAGA	TTGTTTCAGA	AATTAAACTA	GATTGGTCTG	AAGATGAGGA	2820
CAAGGAAGTC	CAAAAGATTG	CCTACAAACG	TCAAATTG	GC	GGAAATTAG GCTTAGATAA	2880

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GTAGAAAAAT AGGGAGTTGG TGAAGATTAT CCTTTACCAAG CGCCCTTTT CTTTTAAAAA	2940
TGAGAAAATT TCGGTATAAT AGTCAAACAA GGTCAAGGTT TAAAGAGAGA GGTGGGTTG	3000
TTATGAGATT TAAAAATACA TCGGATCATA TTGAGGCCTA CATCAAGGCG ATTTAGATC	3060
AATCTGGTAT CGTGGAGTT CAACGGAGTC AGTTGGCAGA TACCTTCAG GTTGTTCCTA	3120
GTCAGATTAA CTACGTGATC AAGACACGCT TTACGGAAAG TAGAGGCTAC TTGGTTGAAA	3180
GTAAGCGTGG TGGCGGAGGC TACATTGTA TAGGACGGAT TGAGTTTCT AGTCATCATG	3240
AAATGCTCCG GGAGCTGCTT TACTCGATTG GTGAGCGAGT CAGTCAAGAA ATTTATGAGG	3300
ATATTCTCCA GCTTTGGTT GAGCAGGAAT TGATGACCAA GCAGGAGATG AATTTGCTAG	3360
AATCAGTAGC TTTGGATCGC GTTTTAGGAG AAGAACGCTC AGTTGTTCGA GCAAACATGC	3420
TACGTCAGAT CATAAAAGAG GTAGATAGAA AAGGAAAGTA AGATGAACTA TTCAAAAGCA	3480
TTGAATGAAT GTATCGAAAG TGCCTACATG GTTGCTGGAC ATTTTGGAGC TCGTTATCTA	3540
GAGTCGTGGC ACTTGTGAT TGCCATGTCT AATCACAGTT ATAGTGTAGC AGGGCAACT	3600
TTAAATGATT ATCCGTATGA GATGGACCGT TTAGAAGAGG TGGCTTGGA ACTGACTGAA	3660
ACGGACTATA GCCAGGATGA AACCTTTACG GAATTGCCGT TCTCCCGTCG TTTGCAGGTT	3720
CTTTTGATG AAGCAGAGTA TGTAGCGTCA GTGGTCCATG CTAAGGTACT AGGGACAGAG	3780
CACGTCCCTCT ATGCGATTTT GCATGATAGC AATGCCTTGG CGACTCGTAT CTTGGAGAGG	3840
GCTGGTTTT CTTATGAAGA CAAGAAAGAT CAGGTCAAGA TTGCTGCTCT TCGTCGAAAT	3900
TTAGAAGAAC GGGCAGGCTG GACTCGTAA GATCTCAAGG CTTTACGCCA ACGCCATCGT	3960
ACAGTAGCTG ACAAGCAAAA TTCTATGGCC AATATGATGG GCATGCCGCA GACTCCTAGT	4020
GGTGGTCTCG AGGATTATAC GCATGATTG ACAGAGCAAG CGCGTTCTGG CAAGTTAGAA	4080
CCAGTCATCG GTCGGGACAA GGAAATCTCA CGTATGATTC AAATCTTGAG CCGGAAGACT	4140
AAGAACAAACC CTGTCTGGT TGGGGATGCT GGTGTCGGGA AAACAGCTCT GGCGCTTGGT	4200
CTTGCCCAGC GTATTGCTAG TGGTGACGTG CCTGCGGAA TGGCTAAGAT GCGCGTGTAA	4260
GAACTTGATT TGATGAATGT CGTTGCAGGG ACACGCTTCC GTGGTACTT TGAAGAACGC	4320
ATGAATAATA TCATCAAGGA TATTGAAGAA GATGGCCAAG TCATCCTCTT TATCGATGAA	4380
CTCCACACCA TCATGGGTTG TGGTAGCGGG ATTGATTGCA CTCTGGATGC GGCAATATC	4440
TTGAAACCAG CCTTGGCGCG TGGAACCTTG AGAACGGTTG GTGCCACTAC TCAGGAAGAA	4500
TATCAAAAC ATATCGAAAA AGATGCGGCA CTTTCTCGTC GTTTCGCTAA AGTGACGATT	4560
GAAGAACCAA GTGTGGCAGA TAGTATGACT ATTTTACAAG GTTTGAAGGC GACTTATGAG	4620
AAACATCACC GTGTACAAAT CACAGATGAA GCGGTTGAAA CAGCGGTTAA GATGGCTCAT	4680

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CGTTATTTAA	CCAGTCGTCA	CTTGCCAGAC	TCTGCTATCG	ATCTCTTCCA	TGAGGGCGCA	4740
GCAACAGTGC	AAAATAAGGC	AAAGCATGTA	AAAGCAGACG	ATTCAAGATT	GAGTCCAGCT	4800
GACAAGGCC	TGATGGATGG	CAAGTGGAAA	CAGGCAAGCCC	AGCTAATCGC	AAAAGAAGAG	4860
GAAGTACCTG	TCTACAAAGA	CTTGGTGACA	GAGTCTGATA	TTTGACCAC	CTTGAGTCGC	4920
TTGTCAGGAA	TCCCAGTCA	AAAACTGACT	CAAACGGATG	CTAAGAAGTA	TTTAAATCTT	4980
GAAGCAGAAC	TCCATAAACG	GGTTATCGGT	CAAGATCAAG	CTGTTTCAAG	CATTAGCCGT	5040
GCCATTGCGC	GCAACCAGTC	AGGGATTGCG	AGTCATAAGC	GTCCGATTGG	TTCCCTTATG	5100
TTCCTAGGGC	CTACAGGTGT	CGGGAAAAGT	GAATTAGCCA	AGGCTCTGGC	AGAAGTTCTT	5160
TTTGACGAGC	AATCAGCCCT	TATCCGCTTT	GATATGAGTG	AGTATATGGA	GAAATTGCA	5220
GCTAGTCGTC	TCAACGGAGC	TCCTCCAGGC	TATGTTAGGAT	ATGAAGAAGG	TGGGGAGTTG	5280
ACAGAGAACG	TTCGCAATAA	ACCCATTCC	GTCTCCTCT	TTGATGAGGT	AGAGAACGCC	5340
CACCCAGATA	TCTTTAATGT	TCTCTTGAGC	GTCTGGATG	ACGGTGTCTT	GACAGATAGC	5400
AAGGGACGCA	AGGTCGATTT	TTCAAATACC	ATTATCATTA	TGACATCGAA	TCTAGGTGCG	5460
ACTGCCCTTC	GTGATGATAA	GAETGTTGGT	TTTGGGCTA	AGGATATTG	TTTGACCAG	5520
GAAAATATGG	AAAAACGCAT	TTTGAAGAA	CTGAAAAAAAG	CTTATAGACC	GGAATTTCATC	5580
AACCGTATTG	ATGAGAACGGT	GGTCTCCAT	AGCCTATCTA	GTGATCATAT	GCAGGAAGTG	5640
GTGAAGATTA	TGGTCAAGCC	TTTAGTGGCA	AGTTTGACTG	AAAAAGGCAT	TGACTTGAAA	5700
TTACAAGCTT	CAGCTCTGAA	ATTGTTAGCA	AATCAAGGAT	ATGACCCAGA	GATGGGAGCT	5760
CGCCCCACTTC	GCAGAACCCCT	GCAACACGAA	GTGGAGGACA	AGTTGGCAGA	ACTTCTCTC	5820
AAGGGAGATT	TAGTGGCAGG	CAGCACACTT	AAGATTGGTG	TCAAAGCAGG	CCAGTTAAAA	5880
TTTGATATTG	CATAAAAGAA	AAAAAGTATC	AGCATCTGAC	CATAAGTCAC	AGTGGAGTGA	5940
AATTCAATGA	AAATCAAAGA	GCAAACACTAG	CAGCTAGCCG	CAGGTTGCTC	AAAACACTGG	6000
TTTGAGGTTG	CAGATAGAGC	TGACGTGGTT	TGAAGAGATT	TTCGAAGAGT	ATGAAACTAA	6060
AACCTATAGC	TTCTAAACGA	TCCGTGGTT	TCATCATTCA	ACACAAAATT	CATATGTTA	6120
TTACCCCTCG	TCGTATTTGT	CTTAGAGCGT	GTGTAGTAGA	AAAAGAGCAG	TCTTATCTGA	6180
AATTTTATT	CTTCAAAAG	AGACCTGTTT	CTTTTTGCA	TGTCAAATCC	GTTCTAGCTG	6240
GTATTTGAAA	AATCAAACCA	ATATTCAATG	AAAATCAAAG	AACAAACTAG	GAAGCTAGCC	6300
GCAGGTTGCT	CAAAACACTG	TTTGAGGTT	GTAGATAGAG	CTGACGTGGT	TTGAAGAGAT	6360
TTTCGAAGAG	TATAAGCTGC	AAGATGAATG	ATTTCCTTGT	ATTGACGTTG	TTGTTGACAA	6420

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AAAGTAGCGG ATAAATGAAA TCCATTCCAT TATCATAGAT GATAGGCTGG TAGGAAATTT	6480
TCAAATAGCA TACAGGAAAT AGATGTATGG AGTTCTGGTA GTAGAAAGGG AGAGAGATGA	6540
ACATTTAGT TGCGAGATGAC GAGGAAATGA TTAGAGAAGG AATTGCAGCA TTTCTGACAG	6600
AAGAGGGTTA TCATGTCATT ATGGCTAAGG ATGGACAAGA GGTCTGGAA AAATTTCAAG	6660
ATCTCCCTAT CCATCTCATG GTACTGGATT TAATGATGCC TAGGAAGAGT GGTTTGAG	6720
TGTTAAAAGA AATCAATCAA AAGCACGATA TTCCTGTCAT CGTCTGAGT GCTCTGGAG	6780
ATGAAACTAC TCAGTCACAG GTATTTGATC TCTATGCTGA TGATCATGTG ACAAAACCTT	6840
TTTCTTGAGT ACTGCTTGTG AAGCGTATTA AGGCCTTAT CAGACGTTAC TACGTCATAG	6900
AGGATCTTG GCGATATCAG GATGTAACAG TGGATTTAC CTCTTACAAA GCACATTATA	6960
AAAATGAAGA AATTGATCTC AAACCAAAGG AATTACTGGT ACTAAAGTGT TTGATTTCAGC	7020
ATAAAATCA AGTTTAAGT AGAGAGCAGA TATTGGAAGA AATTTCAAAAA GATGTAGCTG	7080
ATTTACCTTG TGATAGGGTC GTTGATGTCT ATATTGTCATC TCTTCGCAAA AAATTAGCTT	7140
TAGATTGTAT CGTGACTGTG AAAATGTTG GGTATAAGAT TAGCTTATGAA TAAAAAATCC	7200
TAAATTATTA ACCAAGTCTT TTTTAAGAAG TTTTGCAATT CTAGGTGGTG TTGGTCTAGT	7260
CATTCAATATA GCTATTATT TGACCTTTCC TTTTTATTAT ATTCAACTGG AGGGGGAAAA	7320
GTTTAATGAG AGCGCAAGAG TGTTTACGGA GTATTTAAAG ACTAAGACAT CTGATGAAAT	7380
TCCAAGCTTA CTCCAGTCTT ATTCAAAGTC CTTGACCATA TCTGCTCACC TTAAAAGAGA	7440
TATTGTAGAT AAGCGCTCC CTCTTGTGCA TGACTTGGAT ATTAAAGATG GAAAGCTATC	7500
AAATTATATC GTGATGTTAG ATATGTCGT TAGTACAGCA GATGGTAAAC AGGTAACCGT	7560
GCAATTGTT CACGGGGTGG ATGTCTACAA AGAAGCAAAG AATATTTCG TTTTGATCT	7620
CCCATATACA TTTTGTTA CAATTGCTTT TTCCTTGTT TTTTCTTATT TTTTACTAA	7680
ACGCTTGCTC AATCCTCTT TTTACATTTC AGAAGTGACT AGTAAATGC AAGATTTGGA	7740
TGACAATATT CGTTTGATG AAAGTAGGAA AGATGAAGTT GGTGAAGTTG GAAAACAGAT	7800
TAATGGTATG TATGAGCACT TGTTGAAGGT TATTTATGAG TTGGAAAGTC GTAATGAGCA	7860
AATTGTAAAA TTGCAAAATC AAAAGGTTTC CTTTGTCCGC GGAGCATCAC ATGAGTTGAA	7920
AACCCCTTTA GCCAGTCTTA GAATTATCCT AGAGAATATG CAGCATAATA TTGGAGATTA	7980
CAAAGATCAT CCAAAATATA TTGCAAAGAG TATAAATAAG ATTGACCAGA TGAGCCACTT	8040
ATTAGAAGAA GTACTGGAGT CTTCTAAATT CCAAGAGTGG ACAGAGTGTG GTGAGACCTT	8100
GACTGTTAACG CCAGTTTAG TAGATATTT ATCACGTTAT CAAGAATTAG CTCATTCAAT	8160
AGGTGTTACA ATTGAAAATC AATTGACAGA TGCTACCAGG GTCGTCATGA GTCTTAGGGC	8220

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ATTGGATAAG	GTTTGACAA	ACCTGATTAG	TAATGCAATT	AAATATTCA	AG ATAAAAATGG	8280	
GCGTGTAA	TC	ATATCCGAGC	AAGATGGCTA	TCTCTCTATC	AAAATACAT	GTGCGCTCT	8340
AA	AGTGACCAA	GAACTAGAAC	ATTTATTG	TATATTCTAT	CATTCTCAA	TCGTGACAGA	8400
TAAGGATGAA	AGTTCCGGTT	TGGGTCTT	AATTGTGAAT	AATATT	TTAG	AAAGCTATCA	8460
AATGGATTAT	AGTTTCTCC	CTTATGAACA	CGGTATGGAA	TTTAAGATTA	GCTTGTAGAC	8520	
AGATTAGTTT	TTTATTAAAG	TTCATATAGG	GTAAACATAA	GTGTGTTATT	CTTTGTGTAG	8580	
ATAAAAGAAA	GGATACTAAT	ATGGTATTAG	CGATTATT	AGTAACATTC	TTTATTGAT	8640	
TGATTTTTT	AAAGCGTTCG	ATAGAGAATG	AGAAACGAAT	CCTTAGCAAT	GGCGGGG	8697	

(2) INFORMATION FOR SEQ ID NO: 124:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 4317 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 124:

AACCATA	CAC	ACGGCAAGGC	AAAGCTGACG	CGGTTGAAG	AGATTTCGA	AGAGTATTAG	60
TTGCCTTAA	AGGCATCCAC	CATCGTTGA	AATTCTTCAT	TTGAGAGAGT	AATCCCTTG	120	
CCCATT	TTAG	TATGGTCTGG	ACTCCAAGCA	CGAATATCAA	ACTTGCAGG	GGCACCATTA	180
AAGCTCACAC	GGTTAATT	CTTGGTCCAA	CCTTTTCGT	TTTCAGAAAG	AGTCAACAAG	240	
TGCTCTCGA	TTTCAAATGT	AAATTCTGCC	ATTTCTTCT	CCTTTTTAG	TTTCATTAGT	300	
TTATTCGTA	AA	CTGTAG	ATTTTAGGAA	AATTTATAT	AATATTGATA	TAAAAGAAGG	360
GAGGCCAATA	TGAGACATAA	ATTCCAGCAA	GTCTAAATA	AAATACATGA	TTTTTAAAT	420	
GGATATGACC	AACTGACCA	GACTGAAACC	AACTCCCTTA	CAGCCACTAT	TGAAGAGGCT	480	
ATCCAGAAC	AAACCGCTGT	TCACCTTATC	TTGTCTGAGA	CAAGCTTAC	AGGTGACATC	540	
ATCAAATATG	ATCAGCAAGG	CCAGCAAATT	ATCGTAAAAA	ATTTTCCAA	AAATGTGAGC	600	
CGGATTATCC	GTATAAGCGA	TATTCAACGC	CTGCGATTG	TCCCCTCAAC	TGTCAAACA	660	
GCCCCAAAAA	ATAGATTAA	GAAAGACTGA	GATGTAGTTG	CTTCATCCCA	CTCTTTTTC	720	
TTAGCGAATT	TGTTCAAAAT	GTAAATGAAC	TGCGATATGA	TCTCCATAAC	CACTTCTTTC	780	
CAAGTCACGT	TGTAAACGAT	AGGAAATGTA	GTGTTCTGCA	ATGGTAATGT	AACCTGCGCC	840	
CAATAAACGA	TGTTCAACCA	TAGATTGAAT	CATACTGATA	GTGCGACGTT	CCACCTTGGC	900	

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TTCTTGAAA TCCAAAACCA CCTTCTTAGT GACTTGAGCA AGATTTGAC GCAAATCATC	960
TGTCAAAACA TAAACAGTTT GGGCTGCCCT CAAGATGGCT TGGTAAATCT TATCTGGATT	1020
AAATTTCAGCA ATTTGCCAT TACGTTGAT TACTTGCATA GGTTTCTCCT TTATTCTTG	1080
TTTTCTTGGA TTTCTGCCAG CATTTCCTCT TCTTCTACTG TCAGTTGATA ATGTTCAAGT	1140
AAATCCGGTC TGCGCTCGTA GGTTTCTTT AACTCTCGT ACAATCGCCA CTGACGAATC	1200
TTTTCATGGT GGCCACTCAT CAATACATCT GGACGACCA TGCTCGATA ATCATAGGGA	1260
CGTGTGTACT GAGGATATTC TAAAAGACCT GAAGAAAAAC TATCATCTTGT GTGGCTAGAC	1320
TCCTTGCCAA TCACCTCTGG AATCAGGCGA ACTGTAGCAT CAATCATGGT CATAGCTGCC	1380
AATTCTCCAC CAGTGAGGAC ATAGTCACCT AGGGAAATCT CATCTGTTAC CAAGGTCTTA	1440
ATGCGCTCAT CATAACCCCTC ATAGTGCCCA CAGATAAAGA TTAGCTCTTC CTCTTGAGCC	1500
AAATCTTCAG CATAAGCCTG ATCAAACACTGC TTTCCAGCAG GATCAAGGAG AATAACGCGC	1560
GGATTTTCTT TTTCAATAGC ATCAAAGGAA TCGAAAATAG GTTGTGCTCT GAGCAACATG	1620
CCCTGACCGC CTCCGTAGGG CTCATCATCT ACATGACGGG CCTTTTCAGC ATTTCTCGA	1680
AAATTATGAT ACTGGATATC CAAGAGCCCT TTTTCTCGAG CCTTTCCAAC GATTGAGTGC	1740
TCCAGTGGAG AAAACATCTC TGGAAAGAGG GTTAAAATAT CAATCTTCAT CGTCTAACCC	1800
TTCTAAGATT TCCACATCGA CCCGTTACT TGGAAATATCA ACATTGAGAA CCACTGGTGG	1860
GATATAAGGT AAAAGCAAAT CACGTTGCC TTTTCGTTTG ACCACCCAGA CATCATTAGC	1920
ACCTGGTTGC AGGATTTCCCT TGATGGTTCC AACCAAGCTA TCACCCCTCAT AGACTTCCAA	1980
ACCGATAATC TCGTGTAGT AAAATTCAACC ATCGTCTAGG TCATTCAAAT CTTCCCTCAGC	2040
GACCTTGAGA CTGTATCCCT TGTACTTTTC GATAGTATTG ATATGGTACA TATCTTGAA	2100
TTTAATAATG TCAAAGTTCT TCTGTTACG GTGGCTAGCG ATGGTCACTG TTTGGACAAA	2160
CTGATCTTTT TCATCAAACA AAACCAGCTC AGCTCCTTTT TTAAACCGTT CTTCTGCAAA	2220
ATCCGTCACA GACAAGACTC GCATCTCCCC CTGTAATCCC TGCGTATTAA CGATTTCCC	2280
AACATTAAAG TAGTTCATCT TGTCTCCCTGT AATCTCCTTT TTTCCATCTT ATTCTAACAA	2340
TTCTCGAATA ATAGCCCAA TTTTTCCGA TTCTGACCAT TGAAATAAT GGTGATTCCC	2400
TCCTAAAATG AGTTTAGTAT TGGAAGTCCA ATATTCTGAT TCTCTGTACT CTTTTCTCT	2460
ATAAGGCTGA CAAAAACAA ATACAGGAAT ATGAGCTTCT ATAGATAACAT CCTCAAATC	2520
TTCCCTCAGTA ATCTCTCCAG ATATCTGAAA TTCTGGATCT TGATTTCCA ACTCTAACCC	2580
TTTTTCTTGC ATTAATTCCC AGATTTTTT ATTCTGTTCA GGACTAAATG TTGCTTGAGT	2640
TAAGTTCTTA AAATAAAGTT CAGGACCACA CTCGTCAATC AGCCTCATCT GCTCTTCCAT	2700

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TTCTGGATAA GGATTTCTG AAAAATCAGC AAACATGACT TTTTAGTTG TCGGTTCAAT	2760
TGCTACTAAA GTCTGACGCT TAATTGGTTT CTCGAGTAAT TTGCAAGCTA AAATTCCACT	2820
CCAACATATGT GCACAAAGTA TATATTCAAGA AATTCTTAAT TCTTCAAGTA CTTCATAAAC	2880
CGCATCTGCA AGATTATCTA GATTTTTCC AGCTTGGTCA TGAATCGAC TCCTACCTGT	2940
GTTCGGAAAA TCAATTGTCA AATAACCAAT TGTAGGAGGA GGTTTTCAA GTATAAGTGA	3000
AAAATTTCA TAACTTGGTA GCAAACCTGC TCCGTTAAA CAAACTAGCA CTTTCTTTG	3060
CTTTTGATAA GTAACAGAGA GGCTACCAAT TTCTGTAGAT ACTTCAAACC TCTTCATAAA	3120
GAAATCCACT GATTCTATAT AATGAATTAT TAAAATCCT TATCCTTTAT TTTATCACGT	3180
TCCAAGGATT TTCTCAAGTT GGAGGAAGGG GACAATATCT CTACTTTCCC TTCAATAATC	3240
CTTCCAAATT ATGTTTATGT TGTTAATTAA TGGCTGCGGT TTTGCTTTTC TCAAAGACAG	3300
TCTTGGTAAG GTCAATATGA TTAATAGCTA CGATTGCGAC GGTGTAGTAA ATGATATCAG	3360
CCAGTTCTCT GGCAAGTTCC TCGTTCGAAT CCTATCCCTT CTTTTCGACC AGAGGCCCTA	3420
TTCAAAACCT CGACTACTTC TCCGACTTCC TCCACTAACT TCATAAAGAG ACCTTCATCA	3480
GTCCGAGACT GCTGTTAATG TTGATTAAG TAGTCTTGGA ATTGCCTAAA CGTTCAATCT	3540
TTTATAGTAT ATTGAAACTA GAATAGTACA CCTTTACTTC TAAAACATTG TTAGAAATCG	3600
ATTTGACTGT CCTGATCGAT TTGTCCTGTT CTTGTTTCAT TTTACTATAT CTTCTATTCC	3660
ACACAAAAAA GCGAGACATC CGTCCC GCCC TTCTTATTTT TCGTCAATAA CGATTCTTAC	3720
TTTTTTGTAT TCAGTTGGGA CAGAGTAGAC AACGTTCTT ATCGCAGAAA TAGTGCAGACC	3780
CTTACGACCG ATTACACGAC CCACATCGCT TTGATCAAGA TTCAATGAT ATTCAAAAAA	3840
TTCTGGTGTAA CCCTCAATCT TGATAGTTAA GGCATCTGGT TGTGAAATTAA AGGGTTTCAC	3900
AATCGCAATA ATGAGATTTT CAATCGTATC CATCTGTCAA CCTACTTTAA ACTTATTTG	3960
AAAATTAGA ATCGTGAAT TTTTCAATA CGCCTTCTTT TGAAAGGATG TTACGTACTG	4020
TGTCTGAAGG TTGAGCTCCA TTAGCCAACC ATGCAAGAAC GCGGTCTTCT TTCAAAGTTA	4080
CTTGGTTTTC AGCAACAAGT GGGTTGTAAG TTCAACTGT TTGATGAAA CGTCCGTAC	4140
GTGGTGAACG TGAATCTGCT ACGTTGATAC GGTAGAAAGG TTTTTCTTA GAACCCATAC	4200
GAGTCAAACG GATTTTAACG GCCATTTTA AAGTCTCATT TCTTTAATT TTTATTCGG	4260
TGAAATAGCT GAGCTATTAA GCACATGTTCA TATTATAGCA GATTCTGGC ATGTGTC	4317

(2) INFORMATION FOR SEQ ID NO: 125:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 4881 base pairs

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(B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 125:

AATTTATTTG ACTGGAAATT GTAGAGGGTT CTCGAAATTT CTTGAATGGT TAAAATAAGG	60
ACAAGAGAAA ACATGGATAT CTATATCCTT GTGCCAAAAA AACCACTGCC CTCCCCAGAC	120
CAACCTGAGG AAAGCAGTGA TTCTTATTT AGGAGTTAGG AATGAATACA CGAAATCAAT	180
TTAGCTGATT ATTTTTGTT TTTCAAGAAT TCATCGTATT GTTTTGCAT TTCGTTCAAT	240
ACTTTTCGTT AGGCACCTTC AGATTTCAAT TTTTCCATCA ATTCTGGAAT CGCTTTATCT	300
GGGTCTACAG TACCAGTGT GATAGCTGTA TCAAATTGTT GCATTGTGTT AGCAATAGCT	360
GAGATTCAG ATTTCACATT GTCAGTATTG AAGATAAAC CAAGCGCTGG AGATTCTTA	420
GCTTCTGCCA ATTCTTCCTT AGAATTTCG ATTTGTTGGT CTGTAACGTT TTCGTTGATG	480
TAAAGGATCC AGTTGTTTACCC AGTGTTCAT CCACCCATGT GAGTGTTCG TTTGTAGCCA	540
TCAAGAACGC GAACACGGTT TTCTTACCT TCAATTTTT CCCAGTTCTT GCCTCTGGA	600
CCGTAAACAA GACCGTCAA GAGTTCTGGG TTCTGATTCA AGAGGTTCAA GATTTCCATT	660
GATTTTCCTT TGTTCTTAGA GTTGTGAG ATGACAAAGT TAGCAACTTG TGTTGTTGG	720
TTTTTCTTGA TGAAGTTAGT AATTGGTTG ATTTGGATAT CTTTGTGCG AACACGTGAA	780
AGCAAGCTGT TACCGTAGTC AGCTGGTCCT ACTGTTCTT CACGAACGAA CCAAGTATCT	840
TGTTGAAGGT CAAAGGAAGT ATCGCTGTT GCGACGTCTT TTGGAATGTA GCCAGCTTCA	900
TAGAATTGTT GAAGAGTCTT CAAGTGTCTT TTGAAACGAG GCACCTCGTA ACGGTTTACA	960
ACTTTAGTAG TATGCCCTTC AAGGTCGATA ACGAATGGAA GACCGTTGCG TACTGGTAG	1020
TCAAAATTAT CAGATGGGAT GAAAACCTTA CCAATAGCAA ATGGTACTAC GTCTGGAGCT	1080
TTTTCTTGA TTTGTTCAA GACTGGCTCA AGAGTTTCGT AAGAAGTAAC ACCTGAAATA	1140
TCGATACCAT ATTTAGCAAG GAGAGTTCCG TTGAAGGCAA AGTTTGAGA TGATGCAACG	1200
TTGGCTGCAA CTGGAACAGC GTAAATCTTA CCATTTACAG TATTACCTT GATGTAAGCT	1260
GGGTCAAGTG CTTGTAAAG GTCTTACCT TCTTTTTGT ACAATTCTGT CAAGTCAGCG	1320
TAAGCACCTT TTTGAGCATT TACAATATAG TTATCTGCAA AGGCAATATC ATAGTTTCA	1380
CCAGATGATG TGATAACTGA CATTTCCTTA CCATAGTCAC CCCAGCCAAG GTATTGGATA	1440
TCCAATTGGG CACCAACTTT TTCTTCATG ATTTGTTGG CATTGCTAA CAATTCAATCC	1500
AAGTTGTCTG GTTGTCAACC GATTGGTAC ATTTGATAA CAGGTTGTC ACCTGAATCA	1560

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GCAGCTTTT	TGCTGTTACC	TGTCAAATTT	CCACAAGCAG	CAAGACCTGC	AGCCAGAGCG	1620
ACTACACTAG	CAGATGCAA	AGCATATT	TTCCAGTTT	TCATGATAAA	AACTCCTTT	1680
TTTATTTTA	AACTTATAAA	CAATGTAATG	ATCTTATACT	CAATAAAAAT	CAAAGAGCAA	1740
ACTAGAAAAC	TAGCCGCAGG	CTGCTCAAAG	CACTGCTTG	AGGTTGAGA	TAAGACTGAC	1800
GAAGTCAGTT	ACATATATCT	ACGGCAAGGC	GACGTTGACG	CGGTTGAAT	TTGATTTCG	1860
AAGAGTATTA	ACTTCACACA	AGGGAAGTTG	GGAACTGAGA	AATGTTATTT	CTCAATAAGC	1920
ACTATTCTTT	CACACCACCG	ATAGTCAAAC	CTTTTACAAA	GTAGCGTTGG	AAAAATGGAT	1980
ACAAAATCGC	GATTGGAAGG	GTTGCAACCA	CAACCATGGC	CATACGACCT	GTTCTTTCG	2040
GTAGAGCAAC	TCCCAGTTGA	CCAATCAAGC	CGACCGCTTT	GGCAATGTAG	TCCATATTTT	2100
GTTGGATTTG	CATGAGCAA	TATTGCAATG	GATACAAGTT	GTCACTCTTG	ATGAAAGAA	2160
GGGGCGTTGAA	CCAGTCATTC	CAGAAACCAA	GAGCTGTTAA	GAGCGTGATG	GTTGCGATAC	2220
CTGGTAGTGA	CAATGCCAA	CAGATTTGGA	AGAAAATCCG	GGCCTCACTG	GCACCATCGA	2280
TACGAGCCGA	TTCTAGAATG	GCTTCTGGAA	TGGTCTTCCTT	GAAGAAGGAA	CGCATCAAGA	2340
TGATGTTAAA	TGGTGGAGAGA	AGCATTGGAA	CAATCAAGGC	CCAAACAGTG	TCACCAAGCT	2400
GAAGTACACG	GGTCACCATG	ATATAACCTG	GTACCAAAC	AGCGTTGAAAC	AACATACTGA	2460
GAAGGACGAA	GATGCTAAAG	AATCTGCGAT	ACTTAAAGGT	TGTCCTGAA	ATAGCGTAGG	2520
CATAGGTTGT	TGTGATAAAC	ACATTGTCA	ATGTCCTAAC	TACGGTTACA	AAGACAGAGA	2580
TGAAGAGGGC	TTGTAGGATT	TTATCCTTAA	ACTGTGCCAA	AAACTCAAAA	CCGTCTAACG	2640
CAAATTGGGA	TGGGAAAGAAG	CTATAGCCGT	ATTGGAGGAG	GCTTTCTCG	TCTGTCACTG	2700
AAATAATGAT	AACGAATACA	AAAGGTAGGA	TACAAGAGAG	GGCAATCAA	CCCGAAATGA	2760
TACTGAAGAA	GATATCTGCT	TTCTTACTGA	AGGAGTGAAT	GCCGACATTA	TCAATTTTT	2820
CTTTTTTAAT	TTTCTTTTTT	GCCATATTCT	CCTCCTTCT	AGAACAAAGC	TGAGTTGGAA	2880
TCGACTCGTC	TTGCAAGCAA	GTITGATAGG	ATAACCAGAA	TCAAACCAAC	AACGGATTGG	2940
TAAAGACCGG	CTGCTGCAGC	CATACCGATA	TCTGCTGTCT	GAGTCAAAC	ATTAAGACAA	3000
TATACGTCCA	AAACGTTGGT	TACATTGTAA	AGCTGACCAAG	CATTGTGTGG	GATTTGATAG	3060
AAGAGACCGA	AGTCTGCGCG	GAAGATATT	CCGACTGCAA	GGATGGTCAA	TACAGTTACA	3120
AGCGGAGTCA	ACTGAGGAAT	GGTTACGTTG	CGAATACGTT	GCCACTTGCT	AGCTCCGTCC	3180
ACTGTCGCTG	CTTCGTAGTA	GGTTGGATCA	ATTCCCATGA	TCGTCGCATA	GTACATGACA	3240
CTGCTATATC	CAAAGCCTTT	CCAAATACCT	AGGAAAAGTA	GGAGATAGGG	CCAGATGCC	3300

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AGGTCAGCGT AGAAATTGAC TTCTTGAGA CCAAGACTTT CCAATAGATG ATTGAACACC	3360
CCTTTATCAA TATTTAGGAA GGCATCTGTA AAGAAAATG TGATAACCCA AGACAAGAAG	3420
TAAGGGAAACA ACATAGAAGT TTGAAAAATC TTCACCATTG TCTTAGAACG GAGCTCGCTG	3480
AGGATAATGG CAATCCCTAC AGATACAAC AAACCTAGAA AGATAAAGCC AAGATTGTAG	3540
AGGACAGTAT TTCGTGTGAT AATAAAGGCG TCTCTGAAAC TAAATAAGAA TCTAAAATTA	3600
TCGAGTCCGA CCCATTTACT ATTTATGATA CTATCTATGA ACCATTACT GGTCATGTGG	3660
TAGTCTTGAGGCAACAC GTTCCCAAAT ACTGGAATGT AAAAGAATAG AATCAACCAG	3720
AGTGCCCCCTG GCAAAACCAT CAAGAGAAAG ATCCAGTTGT CTCTCAATGT TTTGAAAAC	3780
TTTTTCATAA TTTCCTCCCT TTTTATTTG ATATCCATCT AAAAATCTT TTTTAGACTT	3840
TTGATAACGA TTACATTATT AGTATACTCC TATTTGCAGG TTAGGTTAAA CTCCTAATTA	3900
TAGAAAAAAAC TCCACAAATT ATGTAGCAGA TTTAAAACCTT TATCACCCT ATCAAACAAA	3960
TGTCCTAAAT CAATTGTTA TTTTATCTCT ATTAGCCCAG TGATGGCGTC ACTCTGTTAT	4020
AAGCATCCAA CAACGGGGTA TACTGAAAAA TCTCCAGACT AGGGAACCTCA GCGATAGTTC	4080
CTAATCTGGA GATTTTAAT ATGTTATTAG GCGTTTGCTT TCAACTTAGC AATAACCTCT	4140
TTAAGATTAT CAATCAACTC TGCTGCAGTA TGCTCAGAGC CTTTTTCATC TGCCAAGAAC	4200
AAAAC TGCTT TTTGAAGTTC TTTTGAGAG TTTTCAAGGA CATCCTATC TACTGTTCA	4260
AGGTTTGAGT CTTAAAGAAG TTTACTTAAT TCCTTGCTA ATTCTTGAG TTTGATTG	4320
AGACTCATCT TCTCCTGCTG TTTCTTGCC CGCTGTTGT CCTCCATCCT TAGTTGCTGA	4380
CTGGCTTCTC TTAATGGACT CTAGGGAAAGC AATGGCATCT TTGACTGTTT GCAAGATATC	4440
ACGTAAACCT TGCTCTGTCA AACTATCATC TGCAAAAGCT TTATTAGCCT CTGCCAAAAC	4500
CAGACGTGCT GAATCTGTGG TAGGATTCGA TACACCTGTC AATGATCTCA AAAGATTTTC	4560
TAAGGTTGAGA GTCTGCTTAC TAATACTAGA CTAAATCAA AAAGTATTAT ATAACAGTGA	4620
TATGAAATCA ACTAAAGAAG AAATCCAAAC CATCAAAACA CTTTTAAAAG ACTCTCGTAC	4680
AGCTAAATAT CATAAACGCC TTCAAATCGT TCTATTTGT CTGATGGCA AATCTTATAA	4740
AGAGATTATA GAACTTTAT AGTAGTTGA AATAAGATGT GAACATCTCT ATCAGGAAAG	4800
TCAAATTAAT TTATAGAAAT ATTTAGCAG CCAAGGTGTA CTGTTATAGA TTCAATACAC	4860
TATACTTGGT GGTTAGCTC G	4881

(2) INFORMATION FOR SEQ ID NO: 126:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13121 base pairs
 - (B) TYPE: nucleic acid

869

(C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 126:

AGGATCCCCG GAAAAGGAGA CTAAAAATGA AGAAAAAAATT TCTAGCATT T	60
TATTCCCAAT TTTCTCATTA GGTATTGCCA AAGCAGAAC GATTAAGATT GTT	120
CCGCCTATGC ACCTTTGAG TTTAAAGATT CAGATCAAAC TTATAAAGGA ATT	180
ACATTATTAA CAAAGTCGCT GAGATTAAG GCTGGAACAT TCAGATGTCC TAT	240
TGACGCAGC AGTCAATGCG GTTCAAGCTG GGCAAGCCGA CGCTATCATG GCAGGGATGA	300
CAAAGACTAA AGAACGTGAA AAAGTCTTCA CCATGTCTGA TACTTACTAT GATA	360
CGTGCATTGC TACTACAAAG TCACACAAAAA TTAGCAAGTA CGACCAATTAA CTGGCAAAA	420
ACGGCTTATC TATTAAAACA TTTGACACTG GTGATTTAAT GAACACAGC TTGAGTGCTG	480
GTGCCATCGA TGCCATGATG GATGACAAAC CTGTTATCGA ATATGCCATT ACCAAGGTC	540
AAGACCTCCA TATTGAAATG GATGGTGAAG CTGTAGGAAG TTTTGCTTTC GGTGTGAAA	600
AAGGAAGTAA ATACGAGCAC CTGGTTACTG AATTAAACCA AGCCTTGCT GAAATGAAA	660
AAGATGGTAG TCTTGATAAA ATTATCAAGA AATGGACTGC TTCATCATCT TCAGCAGTGC	720
CAACTACAAAC TACTCTCGCA GGATTAAGAAG CTATTCCGT TAAGGCTAAA TATATCATTG	780
CCAGCGATTC TTCTTTGCC CCTTTTGTCT TCCAAAATTC AAGCAACCAA TACACTGGTA	840
TTGATATGGA ATTGATTAAG GCAATCGCTA AAGACCAAGG TTTTGAATT GAAATCACCA	900
ACCCCTGGTT TGATGCTGCT ATCAGTGCTG TCCAAGCTGG TCAAGCCGAT GGTATCATCG	960
CTGGTATGTC TGTACAGAT GCTCGTAAGG CAACTTTGA CTTCTCAGAA TCATACTACA	1020
CTGCTAACAC CATTCTGGT GTCAAAGAAT CAAGCAATAT TGCTTCTTAT GAAGATCTAA	1080
AAGGAAAGAC AGTCGGTGT AAAAACGGAA CTGCTTCTCA AACCTTCCTA ACAGAAAATC	1140
AAAGCAAATA CGGCTACAAA ATCAAAACCT TTGCTGATGG TTCTTCAATG TATGACAGTT	1200
TAAACACTGG TGCCATTGAT GCCGTTATGG ATGATGAACC TGTTCTCAA TATTCTATCA	1260
GCCAAGGTCA AAAATTGAAA ACTCCAATCT CTGGAACCTCC AATCGGTGAA ACAGCCTTG	1320
CCGTTAAAAA AGGAGCAAAT CCAGAACTGA TTGAAATGTT CAACAAACGGA CTTGCAAACC	1380
TTAAAGCAA CGGTGAATTC CAAAAGATTC TTGACAAATA CCTAGCTAGC GAATCTCAA	1440
CTGCTTCAAC AAGTACTGTT GACGAAACAA CGCTCTGGGG CTTGCTTCAA AACAACTACA	1500
	1560

870	
AACAACTCCT TAGCGGTCTT GGTATCACTC TTGCTCTAGC TCTTATCTCA TTTGCTATTG	1620
CCATTGTCAT CGGAATTATC TTCGGTATGT TTAGCGTTAG CCCATACAAA TCTCTTCGCG	1680
TCATCTCTGA GATTTCGTT GACGTTATTC GTGGTATTCC ATTGATGATT CTTGCAGCCT	1740
TCATCTCTG GGGAAATTCCA AACTTCATCG AGTCTATCAC AGGCCAACAA AGCCAATTA	1800
ACGACTTTGT AGCTGGAACC ATTGCCCTCT CACTCAATGC GGCTGCTTAT ATCGCTGAAA	1860
TCGTTCTGG TGGTATTTCAG GCCGTTCCAG TTGGCCAAT GGAAGCCAGC CGAAGCTTGG	1920
GTATCTCTTA TGGAAAAACC ATGCGTAAGA TTATCTTGCC ACAAGCAACT AAATTGATGT	1980
TGCCAAACTT TGTCAACCAA TTCGTTATCG CTCTTAAAGA TACAACATAC GTATCTGCTA	2040
TCGGTTGGT TGAACTCTTC CAAACTGGTA AGATTATCAT TGCTCGTAAC TACCAAAGTT	2100
TCAAGATGTA TGCAATCCTT GCTATCTTCT ATCTTGTAAAT TATCACACTT TTGACTAGAC	2160
TAGCGAACAG CTTAGAAAAG AGGATTCGTT AATGGCAAAA TTAAAATTG ATGTAATGA	2220
TTTACACAAG CACTATGGAA AAAATGAAGT CCTAAAAGGA ATTACGACTA AGTTCTATGA	2280
AGGAGATGTT GTTTGTATCA TCGGTCTTC AGGTTCTGGT AAGTCAACTT TCCTCCGTAG	2340
CCTCAATCTT TTAGAAGAAG TCACTAGCGG TCACATCACT GTGAACGGCT ATGATTTAAC	2400
TGAAAAAAACA ACCAATGTTG ACCACGTCCG TGAAAATATC GGCATGGTAT TCCAACACTT	2460
CAACCTCTTC CCTCATATGT CTGTATTGGA CAACATCACC TTTGCTCCTA TTGAGCACAA	2520
GTTGATGACT AAGGAAGAAG CTGAGGAATT GGGATGGAG TTGCTTGAAA AGGTTGGACT	2580
AGCAGATAAA GCTAATGCCA ATCCAGATAG CCTATCAGGT GGTCAAAAAC AACGTGTGGC	2640
CATCGCTCGT GGCCTAGCAA TGAATCCAGA CATCATGCTC TTGATGAAAC CAACTTCTGC	2700
CCTTGACCT GAGATGGTG GAGACGTACT TAACGTTATG AAGGAATTGG CTGAGCAGG	2760
CATGACCATG ATTATCGTAA CCCATGAGAT GGGATTTGCT CGTCAGGTTG CCAACCGCGT	2820
TATCTTTACT GCAGATGGCG AGTCCTTGA AGACGGAACA CCTGACCAAA TCTTGATAA	2880
CCCACAACAC CCTCGTCTGA AAGAGTTCTT AGATAAGGTC TTAAACGTCT AAACCTCAAAC	2940
TGTAAGGATT TCCTTGCAGT TTTTCTACCT CGTATTGGAA TTTTGATTT TTGAAATTT	3000
TATGTTAGAA TTAAGTTAT GAAATGAGGT TTCCTCATAC CTAGCAAGAC TAGGAATAAA	3060
AATAGAAATT AGGTAGCTAG ATGTCATCTA AGGTTATTGT TACAATTTC GGTGCGAGTG	3120
GAGACCTGGC TAAACCGAAG CTCTACCCTT CCCTTTTAG ACTATATCAA TCCGGCAATC	3180
TTTCCAAGCA CTTTGGCCGTT ATTGGAACTG CCCGTAGACC TTGGAGTAAG GAATATTG	3240
AATCTGTAGT TGTCGAGTCC ATCCTTGATT TGGCAGATAG TACCGAGCAA GCCCAAGAAT	3300
TTGCTAGCCA CTTCTACTAT CAAAGCCATG ATGTCATGA TTGGAACAT TATATTGCTT	3360

871

TGCGTCAATT ACAAGCTGAG CTTAATGAAA AATACCAAGC TGAACACAAT AAGCTCTTCT	3420
TCTTGTCTAT GGCACCTCAG TTCTTTGGAA CCATTGCCAA ACACCTCAA TCTGAAAACA	3480
TTGTCGATGG CAAAGGTTTT GAGCGCTTGA TC GTGAAAAA ACCATTGGT ACAGATTACG	3540
CAACTGCAAG CAAGTTGAAT GACGAAC TCC TAGAACATT TGACGAAGAA CAAATTTCC	3600
GTATCGACCA TTATCTTGGT AAGGAAATGA TCCAAAGCAT CTTTGAGTT CGCTTGCAA	3660
ACTTGATTT TGAAAACGTT T GGAACAAGG ATTTTATCGA CAATGTTAA ATTACCTTG	3720
CGGAGCGCTT GGGTAGAA GAACGTGGTG GCTACTATGA CCAATCCGGT GCCCTCCGTG	3780
ACATGGTCCA AAACCACACT CTACAACTTC T T TCGCTCCT CGCCATGGAC AAACCAGCAA	3840
GCTTCACAAA AGACGAGATT CGTGCTGAAA AGATTAAGGT CTTTAAAAAC CTCTATCATC	3900
CAACTGATGA AGAACTCAA GAACACTTTA TCCGTGGCA ATACCGCTCT GGTAAGATTG	3960
ATGGCATGAA ATACATCTCT TATCGTAGCG AGCCAAATGT GAATCCAGAA TCAACAACTG	4020
AAACCTTAC ATCTGGTGCC TTCTTGTAG ACAGCGATCG ATTCCGTGGT GTTCCTTCT	4080
TTTCCGTAC AGGTAAACGA CTGACTGAAA AAGGAAC TCA TGCAACATC GTCTTAAAC	4140
AAATGGATT TATCTTGGA GAACCACTTG CTCCAAATAT TTTGACCATC TATATTCAAC	4200
CAACAGAAGG CTTCTCTCTT AGCCTAAATG GGAAGCAAGT AGGAGAAGAA TTTAACTTGG	4260
CTCCTAACTC ACTTGATTAC CGTACAGATG CGACTGCAAC TGGTGTCT CCAGAACCAT	4320
ACGAAAAATT GATTTATGAT GTCCTAAATA ACAACTCAAC TAACTTTAGC CACTGGATG	4380
AAGTTTGTGC GTCATGGAAG TTGATTGACC GTATTGAAA GCTCTGGCT GAAAATGGTG	4440
CCCCACTTCA TGACTATAAA GCTGGAAGCA TGGGACCTCA AGCCAGCTTT GACCTACTTG	4500
AAAAATTCCGG TGCCAAATGG ACTTGGCAAC CAGATATCAC CTATCGCAA GATGGTCGCT	4560
TAGAATAAAA AAATTCCTG CAAGTTTATG CcTTGCAGGA TTTTTGCTTC TGATTAGATT	4620
AAACCTTCCA AGAGACCTT CATAAAAGTT TCTGAGTTAA ACTCTCCAAT ATCATCGATT	4680
TTTTCACCAA AACCAATCAA TTTTACAGGA ATATTGAGTT CTTCACGAAT GGCTAGAAC	4740
ACACCTCCCTC GAGCAGTTCC ATCAATCTTA GTCAAAACAA TTCCCGTTAA AGGTGTGATT	4800
TTCGAAAATT CTTTGGCCTG TACTAGGGCA TTTTGACCTG TTGATGCATC AAGTGCCAAG	4860
AAGGTTTCAT GTGGTGTCTC TGGCACAA CGTTTGATAA TACGACCAAT CTTTTCCAAC	4920
TCAGCCATAA GGTTATCCTT ATTTGCGAGA CGACCAAGCAG TATCAATCAT GAGAATATCG	4980
ATACCTTCAG TCACGGCACG TTCCATACCA TCAAAGACCA CGCTGGCTGG ATCAGTTTT	5040
TCAGGTCCAG TTACTACTGG AACATCTACT CGTCGGCCCC ATTCAGCTAG CTGAGCTACT	5100

872	
GCACCCGCAC GGAAGGTATC TGCTGCAACC AGCATGACCT TCTTACCAGC TTGTTTGTAG	5160
CGGTGGGCTA GTTTCCGAT AGAAGTTGTT TTCCCAACAC CATTACACACC AACAAAGAGC	5220
ATAACTGTCA AGTTATCTTG GAAGTGGATG CTTTCATCGT AGCTACCATC CTTTCATAA	5280
AGCTCAACCA ATTTCTCAAT GATGACACGA CGAAGTACAT CAGGTTCTT GGCATTTCA	5340
AGCTTGGCTT CGTAACGTAG TTCCTCCGTT AAGTTAGAAG CGACTTGGAC ACCAACATCA	5400
CTCATAATCA GCAGTTCTTC CAGTCCTCG AAAAATTCTT CGTCAACAGA GCGGAAGTTA	5460
GCAAAGAAGG CATTCAAGCG GCCACCGAAA CCTGTGCGAG TTTTCTTAAG ACTGCGGTCA	5520
TATTTTCCTT GAACAGTTTC TTCTGTTGA GGAGCTTCTG GTTCAAGCAC TTCAGAATTA	5580
TTTTCTTCTA CAGTCCTTC GTGCTCAAGC TTCTCTTCCT CTGGTAATTC TTCTGAGTTT	5640
GGTAATTCTT CTATTTCTTC TTGAGAAACC CCTACAGCTG GCTCTGAATC CTGACTTTCT	5700
TCAACTGTGT CTTGGATTTC CTCTCTTGG AACACAGCTT GTTCAACAAT TTCAACCTCT	5760
GCTTCTTCCT GAGAAACTTC CTCAACTTCT GTGAAGGTAG GATCAACATC TTCAGACAAA	5820
TCAAGATTTT CCAGAGCTTC TTTTACAAC TCTTCGATTT TAGGTTCTTC TTTTTTCCG	5880
AATAGACGGT CAAACAATCC CATATCTTAG TTCTCCTTTA GCACATATTC TTCGATAGCC	5940
CAGGCGACAG CTTCCCTCATC GTGGTCATC GGCGTCACTA CATTGCGGC TGCCTTTACT	6000
TCAGGAACAG CGTTTGTCAT AGCAACACCA AGACCTGCC ATTCAATCAT AGAGAGGTCA	6060
TTGGCCTCGT CACCACAAGC CATCACTTGA CTTTGGTCGA TTCCAAGATG GCTGATTAGT	6120
TTTGCCAAAC CTGTTGCTTT ATGAACATTC TTTGGTGACC ATTCTAGCAA CATTTCACGT	6180
GATTTAAAGA TTTCATATTG GTCAAACAAT TCTGGAGAAA TCTTCTGAAT GGCTGCATCC	6240
AAGGGTTCTT GAGCAAAGGC AGTCACGCAT TTGTTGACT TCATTTGACT AGATAAGTCT	6300
TCAAAGTCCA CTGGAACAAA GGTCAAAGCT GGATTGAATT TGGCATAAAG ACTTTCTTGG	6360
TCCGATTGGA TTTGATAAAC TGTTCTTCT GAGATGGCAT CAAGAGGCAG TGATAATTTC	6420
TCTGTTTCTT CATAACAAAC TGCCACATCA TCATATGAAA AGACTGTTTT ATCAAGGATT	6480
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AATACGACCT TGATACCACG ATCACCGCGA gCTTGCAAGG TTTCTTGGT ACGATCCGTC	6660
AGCCTTTAT CAGTAGTCAG CAAGGTCCCG TCCAAGTCCA ATGCAATCAA TTTTATATCT	6720
GCCATTATAA GCCCTCCATA TAAGCTATAA CCGACCGTTC CTTATGGTGA CCAATCACAG	6780
TCTTTGCTAA TTCTAAAATT TCAGGTCGTG CATTTCAGG AGCTACAGGA TGTCCCACAA	6840
CCTGCATCAT ATGTAAGTCA TTAAGATTGT CTCCAAAAGC CATGACCTGA TCCATTGTGA	6900

873

TACCAAGTTT	TTTAACATAAT	TCAACAATGG	CCACTCCCTT	ATCGACATAG	TCCAGAACAA	6960
TATCAATGGA	TTCAAAGCCA	GTTGTCATGG	CCTTAACACC	AGGAACGTTT	TCGTTACCC	7020
AAGCCTCCCC	ATCTTCCAGC	GTTTCTTCTG	TGAAGTTGGT	TGTAAATTG	AAAATGTCAT	7080
CTGTGATATC	TTCCAAACTC	GCTACTTTT	GGATATTTTC	ATTATAGTGC	TGACTCACTT	7140
TCAAATAGGT	CTCATCAACC	GTATCTAGAA	CATATGAACC	CTTCTTACCC	GTCAAGAGCA	7200
GTTTATTGAT	ATCTACATAA	GGTGAAGTTT	TCAGCTTTC	AAAAGTGCC	AGATAAAAGT	7260
CACGAGACAT	AGTCGCTTCA	TACAAGTCCT	GACCTTGATA	CTCTACCAAA	CTGCCATTTC	7320
CCCGCATGAA	AATAATGTCA	TCACGAACAC	CAGCAAATAA	TTTTCTAGA	GACAGAAATC	7380
CCCGACCCGA	AGCTACCGCA	AAGTAAATCC	CTTTTCCCTT	GTAAGGAAACC	AAGAGAGACT	7440
TGAGACGATC	CATATCAAAG	CGTCCATTCC	CATCTAGGAA	GGTTCCGTCC	ATATCCGTTG	7500
CTACTAGTTT	AATTGTCATC	CTTCAATACT	TTCTAAATCT	TTTAACTTAA	CTGAAACAAT	7560
CTTTGAAACA	CCCGATTCTT	GCATGGTCAC	TCCATAGATG	GAATCAGCCG	CTGCCATGGT	7620
TCCCTTACGG	TGGGTTACGA	CGATGAACTG	GCTGCTCTG	TCAAAGCGGT	TGAGGTAATC	7680
CCCAAAACGT	TTAACATTGG	CTTCATCCAG	CGCAGCTTCC	ACCTCATCCA	AGATAACAAA	7740
TGGAATAGTC	TTGACACGAA	TAATGGAGAA	GAGCAAGGCA	AGAGCCGATA	GGGTTTTTC	7800
ACCACCACTC	ATGAGATTAA	GAGACTGGAT	TTTCTTGCC	GGTGGTTGGA	CAGAAATTTC	7860
AACCCCAGCT	GTCAGCAAGT	CTCCTTCAGT	CAAATGAGG	TCAGCCTGAC	CTCCACCAAA	7920
CATCTGCTTG	AAGGTCACTT	TAAAGGACTC	ACGAATGACC	TCAAAGTTG	ATTTAAAGCG	7980
TTCCTTGACC	TCATCATTCA	TCTCTGTAAT	GGTCTCAAGG	AGCAGGTTT	TCGCAGACAA	8040
AATATCATCA	CGTTGGCTAT	TTAGGAAATC	CAGACGGTTG	TGAACTCTT	CGTACTGTT	8100
AATAGCGTCT	AAATTGACAG	GACCAGTGA	GCGTATAGCC	TTCTCTAAAT	CCTTAACCTC	8160
TTGCTCTGCC	AGATTGAGAT	TTTCCAACTC	ATGCGCCTT	TCTAAAGCTT	CTGTGTAGCT	8220
GATCTGGTAC	TGGTCTGTTA	ATTGACTTTG	TAGATGGCGC	AAGCGCTCGC	TAACCTTTTC	8280
TTTCTTGGCT	TCAGCACGAG	TTTGCCTGCG	AATCCACTCT	TCATTCTGCT	GGCGAGCCTG	8340
ATCCAAATGA	CTAGCAATAT	CATCCAGTTG	ACCCTCAATA	TCATCCAAC	CAAAC TGCTT	8400
GCGAATCAAA	CCTTGTGGA	GATTGTTTT	TTGAGTTTTG	GATTCTCCG	CCTGTTGACT	8460
GAGCAATTCT	GTATCAACCT	TCTCAAGATT	ATCAATCTT	TCTTGAAGAA	GGCGCTGGAT	8520
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TTCATAACGT	TTTGCCCTT	GCAGTTCTGT	CTTAAGCAAA	CGAGCTTGCG	CTAGCTCTTC	8640

874	
CTGCAAGTTTGATAGCGTTCTGGATGGCATTGGATGTTAAGCTTAATCTCTTCAATCTC	8700
AGCTTCCAGATTTGCTTGTCACTGGAGATTCGCAGCAAGACGCGCTCTGGCAGTTTCCTT	8760
ATCCGCTTGC CAATCTCCCT CGGAAAGACG ATCTATTTCTCTTCTTGGAGTTCCAAAG	8820
AGTTCCAGTCTTCACATTGCTGACTAGTTGCTGATAAGCAGGAACAAGCCTTGCTC	8880
CTGAATACGTGCCTGCTCTCTTGAGATTTAATAGCTTCTAATGACTCGCTAACATCTGGC	8940
CATCTCATCTTGCAAGGTCTTCAAAGTCGCCTCTCTGAA CCCAAGCTGCTTCTTC	9000
AGCAATTCTTTGTAAATTGCTCCAGTTC TGGCTTGATAAAAATGCTGT TATTCTGGCG	9060
ATTGGCACCA CCTGCATAAGAACCACCTGTGCGCAACTCTGTCCTCCATCCAATGTCACC	9120
ACGAACCTGATAACGAACCTTGGCGAGCTGCTGCACGCGCA TGTTCTACGGTATCAAAGAT	9180
AGCCGTCGTA GCTAGCAAGTTCTGAAAATGGCTTCCAGTCTAGTATCAAAGTCACCAA	9240
CTCATCTGCCATCCCAAGGA AACCTGGGCTTACAGCGATA GCATCTGGTTCTGACTAGA	9300
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TTAAGGAAGTCAATAGCCTTGGTTGCCGA CTCTTCATCTTCTACGATGATATGCTGGCT	9420
ACTTGCCCCCTAAGGCAATCTCTAGGGCACTTGTATAACATCAAAGGTCAGATGCTC	9480
ACTGACTGCA CCAATAATCCCACCTAGCGATCTTTCTCTGGAGAACACCTTTAACACC	9540
TGCATAAAAGTTACTATGATTTCAGGATTTTCCAAA CTGGAGCTCTGGCCTGCTT	9600
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CTTGGCAGTTTCAAGCTCTTCTTTGCTGACTAGCCTCTCTTTAGCTATAGCTAATTG	9780
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CAAACGGGCTTGTGCCTCCTGTTGATTCAA GGCCACTTGC TCGGACTCCA GTTTCGATAG	10140
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TTTGGCCATTTCAGCCTGTAATCTTGGCGTTGCTTTAAGAGTTGATTTCTTCTTC	10260
TAATTTTCA CGCTTTGGTAAATAACTCATCAAGAGTTCTGAAACCTGAGTCAACTCTTC	10320
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AATAGCCTAACGTTGTCCTTCCAAGTCTAAAACTTACGGGCATTCTCAGCTTCTTC	10440

875

AAGAGGCTTG ATTTGATTAT CCAACTCGTA GATAATGTCC TCTAACCGGT CCAGATTATC	10500
CTGAGTTTGC TGCAAGTTAC TCTCGGTTTC TTTTCTGCGA GTCTTGTATT TTAAAACCTCC	10560
AGCAGCTTCT TCACAAATAG CTCGTCGTTC CTCAGGCTTG GAATTAAAAA TCTCCTCAAC	10620
CTTCCCTTGG GAAATAATAG AGAAGGAATC TCGTCCCAAT CCAGTATCCA AGAAGAGGTC	10680
ATGAATATCA CGCAGACGGA CTTCTTGCC GTCAATCTTG TATTCGCTAT CTCCACTACG	10740
ATAGACATGG CGTTCCACCC TGATTTCTTG ACCTGCATCC TTGATAAAATC CGTCATGATT	10800
ATCCAGAGTC ACAACTACAG AAGCATAATT GAGCGGTTTG CGACTTTCGG TTCCAGCAA	10860
GATGATATCC GGCATCTTGC CCCCACGGAG ACTCTTGACA CTAGACTCCC CCAAAGCCC	10920
ACGCAGACTT TCTGTAATAT TGGACTTTCC AGATCCATTG GGTCCAACAA CTGCCGTAC	10980
ACCTTGGTCA AAAACGACCT TGGTCTTATC AGCAAAAGAC TTGAACCCCT GAATTCGAT	11040
TTCCTTTAAA TACATGAATC CAGCCCCTC TCAACGGCAT TTTTGGCAGC TTCTGCTCT	11100
GCTAATTTCT TAGAACGACC TTGGCCTTGA CGCATGCTCT TACCTTCAAC AAGAACTTCT	11160
ACATCAAAAA CCTTATCGTG AGCAGGCCCT GTTTCAGAAA TCACCTGATA ACGAATAGCC	11220
ACATCACCAC TGACCTGAAG CAACTCTTGG AGATGGGTTT TATAGTCTGT AATCATCTCA	11280
AACTCGCCTG CTTCAACCTT AGGAATCATG ACTTGATAGA TAAATCCCTT GACCTTGGCC	11340
ACATCCTTAT CCAAAAGAAG GGCACCAAGA AAGGCTTCAA AGGCATCACC AAGAATGGTG	11400
TCACGATTGC GACCACCTGA TTTTCTTCC CCTTTACCCA ACTTGATAAA CTGGTCAAAC	11460
TGGCAATCAC GCGCAAAACC AGCTAAACTC TCCTCACGGA CAATCATAGC ACGGAGTTT	11520
GATAGGTAC CTTCAGGCTT TTTAGGATAT TTTTTATATA GATATTCTGA AATCAATAAC	11580
TGTAGAACAG CGTCTCCTAA AAATTCCAAG CGTTCATTGT GTGAAATTAA TAAGAGGCGG	11640
TGCTCATTGG CATAACTCGT ATGAGTAAAG GCAGTTCCA GTAACTTTT GTCTGAAAT	11700
TCGATTGCAA AATGATTCTT TAGTACAGTT TGTAATTCTT TCATACCAAC CTCTTCTAA	11760
CTGATAATAG TCCTTTTAT TATATCAAAA AAAGCCCCCT GAGTCACTCT AAAACGGGAC	11820
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AGGATAAAGA AAAAAGCCCT ATTAAAGGCT TTTTAGGATG TTTACATCCA CCCTGAGGGA	11940
ATCGAACCCC CATCTCAAGA ACCGGAATCT TACGTGATAT CCATTACACT AAGGGTGGAA	12000
ACTTGTTTTA TTATAACAGA AATTGCTCT AATAACAAGT TTTTGGTCA AAGACCCGT	12060
CTTAGTGGGA AGCATCCCCA TTCCAGATGG AGTTTTCAC GATCACATAA TCAACGTGTT	12120
TAAGGTCAGC AACCTGACGT CCACCTGCAT AAGAAATAGC ACTTTGAAGG TCTTGTCCA	12180

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TCTCAGTTAA	AGTGTCTTGC	AGATGACCTT	TAGCAGGAAG	CAAGATACT	TTGCCTTCCA	12240
CATTTTTGTA	AGCACCTTTT	TGATATTGTG	AGGCTGAACC	ATAATATTCT	TTGAACTGTT	12300
CACCATCGAC	TTCAATCGTT	TTCCCTGGAC	TTTCAATGTG	TCCTGCAAAG	AGGGAACCAA	12360
TCATGATCAT	GCTAGCACCG	AAGCGGATAG	ACTTAGCAAT	ATCACCGTGA	GTACGAATT	12420
CTCCATCAGC	GATAATCGGT	TTACGCGCAG	CCTTGGCACA	CCAGCGTAGA	GCAGCCA	12480
GCCAACCACC	TGTACCAAAA	CCAGTCTTAA	CCTTGGTGAT	ACAAACCTTA	CCAGGACCGA	12540
TTCCGACCTT	AGTAGCATCC	GCACCAGCAT	TTTCCAATT	ACGCACAGCT	TCTGGTGTTC	12600
CCACATTTCC	AGCAATGACA	AAGGTATCTG	GCAATTCTT	CTTGATGTGT	TGAATCATAG	12660
AAATCACGCT	ATCCGCATGA	CCATGAGCAA	TATCAATAGT	GATATACTCA	GGAGTATCAG	12720
CCTTGAGCTG	GCTAACAAAA	TCATACTCAT	AATCCTTAAC	ACCGACAGAG	ATAGAACCAA	12780
TGAGGCCCTTG	ATTGTGCATT	CGTTAATAA	AAGGAATGCG	TCCTGCCTCA	TCAAAACGGT	12840
GCATAATGTA	GAAGTAACCA	CCTTAGCCA	GTTGCTCTGC	TACATTTCA	TCCAAAATCG	12900
TCTGCATATT	CGCTGGCACA	ACAGGTAGTT	TAAAGGTGTG	ATTTCCCTAA	GTGACACTTG	12960
TATCCGCTTC	TGCACGGCTT	TTAATGACAC	ATTATTTGG	AATCAATTGA	ATATCTTCGT	13020
AATCAAAAT	TGGAAATTCA	TTAACACATAT	CGATGTCTCG	TTTCTTTGT	AATGACCTAC	13080
CTATGCTCTT	GCATCACTAC	GCCTTTCCG	ACGTTTCCTG	G		13121

(2) INFORMATION FOR SEQ ID NO: 127:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9578 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 127:

CCGAATGCAA	TGTTTACGGT	TGAACCTTGAA	AATGGACATC	AGATTTAGC	AACAGTTCT	60
GGTAAAATTC	GTAAAACTA	TATTCGTATT	TTAGCGGGAG	ATCGTGTAC	TGTCGAAATG	120
AGTCCATATG	ACTTGACACG	TGGACGTATC	ACTTACCGCT	TTAAATAATC	AAAAAACTTG	180
GAGGGATAAG	AAATGAAAGT	AAGACCATCG	GTCAAACCAA	TTTGCATA	CTGTAAAGTT	240
ATTCGTCGTA	ATGGTCGTGT	TATGTAATT	TGCCAGCAA	ATCCAAAACA	CAAACAAACGT	300
CAAGGATAAG	ATAGAAAGGA	GAAAACATGG	CTCGTATTGC	TGGAGTTGAT	ATTCCAAATG	360
ACAAACGCGT	AGTAATCTCA	TTGACTTATG	TTTATGGTAT	CGGACTTGCA	ACATCTAAGA	420
AAATTGGC	TGCTGCTGGA	ATCTCAGAAG	ATGTTCGTGT	ACGTGATCTT	ACATCAGATC	480

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AAGAAGATGC TATCCGTCGT GAAGTGGATG CAATCAAAGT TGAAGGTGAC CTTCGTCGTG	540
AAGTAAACTT GAACATCAAA CGTTGATGG AAATCGGTTC ATACCGTGGT ATCCGTCACC	600
GTCGTGGACT TCCTGTCCGT GGACAAAACA CTAAAAAACAA CGCCCGCACT CGTAAAGGTA	660
AAGCTGTTGC GATTGCTGGT AAGAAAAAAT AATATAGGAG GTAAAAGTCT TGGCTAAACC	720
AACACGTAAA CGTCGTGTGA AAAAGAATAT CGAATCTGGT ATTGCTCATA TTCACGCTAC	780
ATTTAATAAC ACTATTGTTA TGATTACTGA TGTGCATGGT AATGCAATTG CTTGGTCATC	840
AGCTGGTGCT CTTGGTTCA AAGGTTCTCG TAAATCTACA CCATTCGCTG CTCAAATGGC	900
TTCTGAAGCT GCTGCTAAAT CTGCACAAGA ACACGGCTT AAATCAGTTG AAGTTACTGT	960
AAAAGGTCCA GGTCTGGTC GTGAGTCAGC TATTCGTGCG CTTGCTGCCG CTGGTCTTGA	1020
AGTAACAGCA ATTCTGTGATG TGACTCCAGT GCCCACACAAT GGTGCTCGTC CTCCAAAACG	1080
TCGCCGTGTA TAATCATCGC ATTACACTGC TTTTCGTTA AGAGGGAGTA ACTAAATGAT	1140
CGAGTTGAA AAACCAAATA TAACAAAAT TGATGAAAAT AAAGATTATG GCAAGTTGT	1200
AATCGAACCA CTTGAACGTG GCTACGGTAC AACTCTGGT AACTCTCTTC GTCGTGTACT	1260
TCTAGCTTCT CTACCAGGAG CAGCTGTGAC ATCTATCAAC ATTGATGGTG TGTTACATGA	1320
GTTTGACACA GTTCCAGGTG TTCGTGAAGA CGTGATGCAA ATCATTCTGA ACATTAAGG	1380
AATTGGCAGTG AAATCGTACG TTGAAGACGA AAAATCATC GAACTGGATG TTGAAGGTCC	1440
TGCTGAAGTA ACAGCTGGTG ACATTTGAC AGATAGCGAT ATTGAAATTG TAAATCCAGA	1500
TCATTATCTC TTTACAATCG GTGAAGGTTC TTCTCTAAAA GCGACTATGA CTGTTAACAG	1560
TGGTCGTGGA TATGTACCTG CTGATGAAAA TAAAAGGAT AATGCACCAAG TTGGAACACT	1620
TGCTGTAGAT TCTATTCTATA CACCAGTTAC AAAAGTCAAC TATCAAGTGG AACCTGCTCG	1680
TGTAGGTAGC AATGATGGTT TCGACAAATT ACCCTTGAA ATCTTGACAA ATGGAACAAT	1740
TATTCCAGAA GATGCTTAG GGCTTCAGC ACGTATTTG ACAGAACATC TTGATTTGTT	1800
TACAAATCTT ACTGAGATTG CTAAGTCAAC TGAAGTGATG AAAGAAGCTG ATACTGAATC	1860
TGACGACCGT ATTTTAGATC GTACGATTGA CGAACTGGAC TTGTCTGTGC GTTCATACAA	1920
CTGTTAAAAA CGTGCCTGTA TCAATACTGT GCATGATTTG ACAGAAAAT CTGAAGCAGA	1980
GATGATGAAA GTACGAAATC TTGGACGCAA GAGTTGGAA GAAGTGAAC TCAAACTCAT	2040
TGATTTGGGT CTTGGATTAA AAGATAAATA AAGGAGGAAT ACATGGCTTA CCGTAAACTA	2100
GGACGCACTA GCTCACAACG TAAAGCAATG CTTCGCGATT TGACAACTGA CCTTTGATC	2160
AACGAATCAA TCGTGACAAC TGAAGCTCGT GCTAAAGAAA TCCGTAAAAC TGTTGAAAAA	2220

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ATGATTACTC TAGGTAAACG TGGTGATTTG CATGCACGTC GTCAAGCAGC TGCTTTCGTA	2280
CGTAATGAAA TCGCATCTGA AAACTATGAT GAAGCAACTG ATAAGTACAC TTCTACTACA	2340
GCACCTCAA AATTGTTCTC AGAAATCGCA CCTCGTTATG CTGAACGTAA CGGTGGATAC	2400
ACTCGTATCC TTAAAACCTGA ATCACGTCGT GGTGATGCA CGCCAATGGC GATCATCGAA	2460
TTAGTATAAA ATCATCAATT TTGTTGAGTG TTATGATGAT GGAGTCTTGT GCTCTTAGTC	2520
TAGCTCTGGT CTACCGCTAG GATTTCGGTC CTAGCGGGAA CACTCATCAT AAGTTGGGAT	2580
AGTAGACGCT TGTTCACGAA ATTGTTTTTT TCTTAAGAAC AACTTCGTAA GCAGGCGTTT	2640
TTGAGTATTT TCGTTAGAAT TATGCTATAC TATTTGAAAA GAATCCTGTT TAATGTTAAG	2700
GTTTCTTATT TTAAGAAGAA TTGGAGTTA CTTATGAAAG CCATTATAAC TGTTGTTGGT	2760
AAAGATAAAAT CTGGAATTGT TGCAGGTGTT TCTGGTAAAA TTGCAGAATT AGGATTGAAT	2820
ATTGACGATA TCTCTCAAAC TGTCTTGGAT GAATATTTA CGATGATGGC TGTTGTATCT	2880
AGTGATGAAA AGCAAGATTT TACCTATCTT CGTAATGAAT TTGAAGCTTT TGGGCAAAC	2940
TTGAATGTA AAATCAATAT TCAGAGTGCA GCGATTTTCG AAGCTATGTA TAATATCTAG	3000
GAGGTCATCA TGGATATTAG ACAAGTTACT GAAACCATCG CCATGATTGA CGAGCAAAAC	3060
TTCGATATTA GAACCATTAC CATGGGGATT TCTCTTTGG ACTGTATCGA TCCAGATATC	3120
AATCGTGCTG CGGAGAAAAT CTATCAAAAA ATTACGACAA AGGCGGCTAA TTTAGTAGCT	3180
GTTGGTGATG AAATTGCGGC TGAGTTGGGA ATTCCTATCG TTAATAAGCG TGTATCGGT	3240
ACACCTATTCT CTCTGATTGG GGCAGCGACA GATGCGACGG ACTACGTGGT TCTGGCAAAA	3300
GCGCTTGATA AGGCTGCGAA AGAGATTGGT GTGGACTTTA TTGGTGGTTT TTCTGCCCTTA	3360
GTACAAAAAG GTTATCAAAA GGGAGATGAG ATTCTCATCA ATTCCATTCC TCGCGCTTG	3420
GCTGAGACGG ATAAGGTCTG CTCGTCAGTC AATATCGGCT CAACCAAGTC TGGTATTAAT	3480
ATGACGGCTG TGGCAGATAT GGGACGAATT ATCAAGGAAA CAGCAAATCT TTCAGATATG	3540
GGAGTGGCCA AGTTGGTTGT ATTCGCTAAT GCTGTTGAGG ACAATCCATT TATGGCGGGT	3600
GCCTTTCATG GTGTTGGGA ACCAGATGTT ATCATCAATG TCGGAGTTTC TGGCCTGGT	3660
GTTGTGAAAC GTGCTTGGGA AAAAGTTCGT GGACAGAGCT TTGATGTAGT AGCCGAAACA	3720
GTAAAGAAAA CTGCCTTAA AATCACTCGT ATCGGTCAAT TGGTTGGTCA AATGCCAGT	3780
GAGAGACTGG GTGTGGAGTT TGGTATTGTG GACTTGAGTT TGGCACCAAC CCCTGCGGTT	3840
GGAGACTCTG TGGCACGTGT CCTTGAGGAA ATGGGGCTAG AAACAGTTGG CACGCATGGA	3900
ACGACGGCTG CCTTGGCCCT CTTGAACGAC CAAGTTAAAA AGGGTGGAGT GATGGCCTGC	3960
AACCAAGTCG GTGGTTTATC TGGTGCCTTT ATCCCTGTTT CTGAGGATGA AGGAATGATT	4020

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GCTGCAGTGC AAAATGGCTC TCTTAATTAA GAAAAACTAG AAGCTATGAC GGCTATCTGT	4080
TCTGTTGGAT TGGATATGAT TGCCATCCCA GAAGATAACGC CTGCTGAAAC TATTGCGGCT	4140
ATGATTGCGG ATGAAGCAGC AATCGGTGTT ATCAACATGA AAACAACAGC TGTCGTATC	4200
ATTCCCAAAG GAAAAGAAGG CGATATGATT GAGTTTGGTG GTCTATTAGG AACTGCACCC	4260
GTTATGAAGG TTAATGGGC TTCGTCTGTC GACTTCATCT CTCGCGGTGG ACAAACTCCA	4320
GCACCAATTTC ATAGTTTAA AAATAAGAA AATAGGAGAA ATTTTAAGTT CTATTAAAGA	4380
TTAGACGTGT ATACTATAAT CATTAAATAA AGACCTCCTA ATATTATTG AAACAGATAA	4440
CACTGAATTA GTTTGAATTG GATTTTCATC TAATATCTT ATTTAATGAA CTCCTAAACT	4500
TTTTCATAAT AATCTCCTTC AAAAGTCGCC TGTATGGGTG GCTTTTATTT TATCATTCTAT	4560
GATATAATAG AAGCAAACGG AGGACGGAAA ATGGTAAAAG TACGATTGTA TTTGGTACGT	4620
CATGGCAAGA CCATGTTAA CACGATTGGT CGCGCGCAAG GTTGGACGAG TACTCCCTTA	4680
ACTGCTGAAG GTGAACGAGG GATTCAAGAG TTAGGAATCG GTTTGCGAGA ATCTGATCTA	4740
CAGTTGAGC GTGCTTATTG GAGTGTATTCT GGTCGTACCA TTCAGACCAT GGGATTATC	4800
CTTGAAGAAC TTGGCTTGCA GGGGAAATC CCTTATCGCA TGGACAAGCG TATCAGAGAA	4860
TGGTGTTCG GTAGTTTGA TGGAGCCTAT GATGGCGATC TTTTCATGGG CATTATTCT	4920
CGTATCTTTA ATGTGGACCA CGTTCACCAA TTGTCTTATG CTGAACCTGGC TGAGGGCTTG	4980
GTAGAGGTGCG ATACAGCTGG TTGGGCTGAA GGCTGGAAA AACTCAGTGG CGGAATCAAG	5040
GAAGGCTTTG AAATGATTGC AAAAGAAATG GAAGATCAAG GTGGAGGTAA CGCCCTTGT	5100
GTCAGCCATG GAATGACTAT TGGAAACCATT GTTTATCTGA TTAATGGCAT GCATCCGCAT	5160
GGTCTGGATA ATGGTAGCGT GACAATCCTT GAATATGAGG ACGGCCAGTT TAGGGTTGAA	5220
GTGTCGGTG ACCGTAGTTA CCGAGAGCTA GGACGTGAGA AGATGGAAGA AGGCTCTATT	5280
TAATCAGTCT AGACTGCTT GCCATGAGCT AGGGATTGTA TAAGAATATC AAGATAAGAA	5340
AAAACAGCCG AGGGCACTCC TTTCGGCTGT TTTTGATGTG GAAAACCTAAA GTGTAATGCT	5400
ATTGCTTTTA GAGATTTCA TAAACAAGAG CAAGGAACCT ACTGTTAGAA CAGTCAGGAT	5460
AGTTGACAAG GTTGCGCTA CACCGTAATT TCCTCTGAGA ACCTCTGTAT AAATAGCTAC	5520
AGTCATTGTT CTTGTTTGA CATTGTAGAG GAGGATAGAA GTAGAGAGTT TTGAAATCAT	5580
TGTGACTCAA GATAAGATGG CTCCAGAAAT GATACCAGAT AGCATCATTG GAGTTGTAAT	5640
CTTAGCAAAG GTATTGAGAC GACTACTTCC TAAGCTTCA GCAGCTTCTT CAATACTTGG	5700
TGCTATTGTT TGTAAGCTAG CAACAGATGA GCGAATAGTA TAAGGTAATC TTCTGGCAGA	5760

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TAGAGACATA ATCAAGATGA AAGCAGTCCC TGTAATCATA AGAAATCCAC TTCCAAATAG	5820
ACCACTATTG AAGGAAGAAA TGAAGGCAAT CCCTAGAACG GTTCCTGGTA CAATATAAGG	5880
TACCATACTG AGGCTGTCAA TTAAGTTGT AAACAAATTG CGTTTCTAA CGGCTAGGTA	5940
GGAGATAAAT GTCGCAAATA GAACAACCTAG AACTAAGGC ATCAAAGGG TACGAATGGT	6000
ATTGAAAATA GCAGATCCC TACGATGGAA AGCTACCTTG TAACTGTTG GAGAATAACC	6060
TTTAACAGAT ACCATACCTG ATGTTTTAG GAAAGAGGTA TAAATTAAGT AGATTTGAGG	6120
TAAAACAGAG ATAAAGATAA TTCCGTAGAC TGTTGCATAA ATGGCAGCCA TTTTCCTT	6180
TGTAGTTTT TTAGGCTCAA TTGGATGGAG CAGATTCATG CTGAAACTGT AGCGGTTGC	6240
AATGTGTTT TGGATAAGGA AAATTGCCAA GGCAATGATA ATCGCCATAA TTGCAAAAGC	6300
AGAATTCCT CCAACCTCGC TAATAAATTG GGTATAAATC AGGACAGGG AAGTCCGATA	6360
CCCTTCGCCA ATCAACATAG GCGTTCCAAA GTCTGAGAAT GCTCTCATAA ATACAAGCAA	6420
GGAGCTGCTA GTAAGGTTGG AACTAGGAGA GGTAAAACAA CCGTTACGAT AGGTTAAAT	6480
CCGAAGGACC CCATGTTTC AGCTGTTCA AGTAGAGAAT TGTCAATACT GTTCATTGTT	6540
CCAGCAACAT ATAGAAATAC CAGTGGGAAT AGTTGCAGTG TAAAGACAAG TACAATTCT	6600
TTGAATCAAT AAATATCGAT AGCTGGAAGA TAAAGGGCAT TTGTAAAAAA TTTAGTGATG	6660
ACCTCATTTC GTCCTAGCAA GAGAACCCAG GAGTAGGCTC CTACGAAAGG AGCTGACATG	6720
GAAGCAATGA TAATCAATAT TTGTAGAAAT TTCTTCCCCCT TGAAGTCATA CATAGAGAAG	6780
AGATAAGCTA ATAGGGTCC TACAACTAAG GAAGTGATAG TAGCGGTAAT GGAAACCTTG	6840
AAACTGTTGA CTAGTGTCTC AGAGTAGTAG GCTTTACTAA AGAAAGTGAC AAAATTAGCT	6900
AGTGAGAATT GTCCTTCATG TATAAGTGCT TGCTTGAGCA CGGTAACGAT AGGATAAACG	6960
AGAAAGATAG GATAGGTAAG AAAGAGGAAG AAAGAGGAAA CTGTCCAAAT ATTTAGTTT	7020
TTACGTTCCA TGGTTGACTC CTTTTATCAG GTTTGGGAA CCATCTGCAG AAAAGATGTT	7080
TAATTTTGTC GTATTGATTC GTAGACGAAT ACGATTGCCT TTTTGTAGAT CTTCTTCAAA	7140
AGTTGATTCT TCACTAACTT GAATTTTGAGA GGCAAAACCT GTCTCAATGA AATAATCCGT	7200
ATTTAGTCCA AGATAGACGC TATCTCTAAT AGTTCCCTCA ATATCTCCAG ATTCACTTT	7260
GATAAACTCT TCGGGACGAA TGCTTACATG AATAGCTTGC TCCTCAACCT GATCAAGAGC	7320
TGGCATTCGA AGGGCATAGC CATCTGAAAA GACGATATAA GCGCCGTCGC TCCGTTTTC	7380
AAGATTGGCA GGGATAATAT TTGTGCGTCC GATAAAGGTT GCCACAAACT CATTAGCTGG	7440
TTTATGATAG AGTTCTTTG GTCGGCCGAT TTGTTGGATC ACCCCATCTT TCATAACAGC	7500
AATTGGTCT GAAATAGCCA TGGCTTCTTC TTGGTCGTGG GTTACATAAA CAGTTGTAAT	7560

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TCCCCACTTCG	TGTTGGATTT	CTCGGATGGC	TTGACGCATA	TCCAAGCGAA	GTTTGGCCTC	7620
CAGATTACTA	AGTGGCTCGT	CCATGAGGAG	AACACTTGGA	TTAACCGCTA	AGGCGCATGC	7680
CAAGGTGACA	CGTTGTTGTT	GTCCACCAC	GAGTTTATCG	GGCTTCGAT	CCGCATATTG	7740
AGCAATTGC	ATGAGTTCAA	GATACTTGTT	GGTCTGTTGA	ATCAATTCTT	CTTTTGGAAC	7800
CTTCTTTGC	ATAAGACCAA	AAGCAACGTT	GTCTCGGACA	GTCAAATGTG	GGAAAATAGC	7860
GTAGTTTG	AAAACCATCC	CGATATTGCG	TTTGCCTGGGT	TCCATATTAT	TGATTTTGT	7920
ATCATCGAAG	TAAAATTCTC	CACCTTCGAT	ACTGTTGAAA	CCTGCAATCA	TACGAAGAAG	7980
GGTCGTTTC	CCACATCCTG	AAGCTCCAAG	AAGGGTAAAG	AGACTTCCTT	TTGGAATTGT	8040
AATGTTCAA	TTCTCAATAA	CAGGGACATC	GTGGTAGATT	TTTTTGGCGT	TAATAATT	8100
GATCTCACTC	ATAGTGAACC	TCTTTTACTG	TTTAGATTGG	ATATCTGTA	AGACTTCGTT	8160
GTATTTCTTA	ACGATATCTG	ATTTATTCTT	GATGACATAA	TCATAATCTT	CAGTGAGTGT	8220
TTTGATTTTG	TCAATTGGTT	TCATGTTTTC	GCTTGTGTTA	GCATTTTAC	GAACAGGACG	8280
GTAGTAGTG	GTTGTACCAA	GTGTATCTTG	TACTTCTTGA	GAGATAATAA	AATCGATAAA	8340
TTTCTGGCA	TTTCCATAT	TTTAGATTT	TTAACGATA	GCAGCACTAG	CAGGTAGGAA	8400
GACGGTCCCT	TCTTTGGAT	AGACTACCTT	AATGTTAGCT	CCGTCATTTA	AGAGTTAAC	8460
TGCTGGATCT	TCATAAGAGA	GACCAACAGC	CATTTCTCCA	TCAGCGACTA	CTTTATAGAC	8520
ACTAGATGAA	CTTGAACCGA	TTTACCATC	AATAAGTGTG	AAAAGATCTT	TTACATAAGA	8580
CCAAGCCTTA	TCATCTTGT	AACCACCTTG	AGCTTGTAGC	ATATTGTTA	ATTGAGCAA	8640
GGCGCTAGAA	GAGTTGCTG	GGTCAGCAGT	TGCGATTTT	CCTTTAGTT	CAGGTTGAA	8700
AAGATCGTTA	TATCCTTCGA	TGTTCATGCC	TTAGTTAAA	TCAGGGTGA	CGATTAAC	8760
ACTACCACCT	AGTGTATAAG	GAGTAGAGTA	GCCAGTTGTG	TTTGATATT	CTTGATAAC	8820
ATTATCATT	TCTTTGAAG	TATAGTTTC	AAAGAGTTCT	CCGTGGTAG	TATATTGTGT	8880
ATAAGAACCA	CCAAAGATAA	CATCAGCTAC	AGGAACCTCT	TTTCTGACT	CTAGTTTT	8940
GAAAAGTTCT	CCAGTACCA	CTTGAATCAG	TTCTACTTTG	ATACCATATT	TTCTTCAA	9000
GGCAGGAATA	GTTGCTCCAA	TTAACGCCCTC	TGAGTTGGT	GAATAAACGA	CTAGCGAAC	9060
GCCGTCTCC	TTATCAGATG	AACTGTCATC	GGCAGATTCA	TTAGAAGAAC	AAGCAGCATA	9120
ATACATCCAT	TTCTTTTCA	TGATGGATAC	CTCCGTTGTG	TTATTTAAGT	TTATTTAAA	9180
ACAATGTAAG	CGTTTTAAA	ACATACAATT	CTATTCTATA	GTGTATTGAA	TCTATAACAG	9240
TACACTTTGA	CTGCTAAAT	ATTTCTATAA	ATTAATTGAA	CTTTCCTGAT	AGAGATGTT	9300

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ACATCTTATT TCAATTCACT ATATTAGAGT AAAATTCTCT ACAAAAAGAA GAATAGCCTA	9360
TTTTACTATT CTTCTGAGTG ATTTCAATTC CTTTGGGGAA ATATGGAGAT ACTTTTAA	9420
TCCTGACAAA TGGTTGTTTC TTTTCTAAA TCGGTGATAC TGTATCGGAG AATGCGCGTG	9480
AGGTACACAAA GGCTGCGATA GAGCTTCTAT GGAGAATTTC TTTTGGAGA GATTTTTAA	9540
AGGAATGAGA CATCCGCTAC CTCCTGGAA GTTTTTG	9578

(2) INFORMATION FOR SEQ ID NO: 128:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 13440 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 128:

CGGGCTGTTG TGACGATTCT TATTCTATC TGTGTTATCT TTTTGGGAAC TATTTTGGGT	60
GTTGCTTGG CTTTTGGCA ACGTTCAAAG TTTAACCGCG TTGTTTGGTT GGCCAACTTG	120
TACGTTTGGA TTTTCCGTGG GACACCGATG ATGGTTCAAA TTATGATTGC CTTTGTCTT	180
ATGCATATCA ATGCTCCGAC TATTCAAGATT GGAATTTCAG GTGTTGATT TTTCGCGTCTG	240
ATTCCAGGGA TTTTGATTAT CTCTATGAAT AGTGGTGCTT ATGTTTCGGA GACTGTTCGT	300
GCCGGAATCA ATGCGGTTCC AAAAGGTCAG CTAGAACGGG CTTATTGCT AGGGATTGCGT	360
CCTAAAAATG CGATGCGTTA TGTGATTTG CCACAAGCAG TCAAAATAT CTTGCCAGCA	420
TTGGGGAACG AATTATCAC CATTATCAAG GACAGCTCCC TCTTATCAGC TATTGGGTC	480
ATGGAGTTGT GGAATGGGC TACAAACAGTT TCTACAACAA CCTATCTACC TTTAACACCA	540
CTTTTATTG CAGCATTTA CTACTTGATT ATGACCTCTA TTCTGACAGT AGCCTTGAAA	600
GCTTTTGAAA AACATATGGG ACAAGGAGAT AAGAAATAAT GACAGAAACC TTGATAAAAAA	660
TTGAAAATTT ACATAAATCC TTTGGAAAGA ATGAAGTATT GAAGGGCATC AACCTCGAGA	720
TTAAAAGAGG AGAAGTTGTC GTTATCATCG GTCCTTCAGG GAGCGGGAAA TCTACCTTGC	780
TTCGCTCTAT GAATTGTTG GAAGAAGCAA CCAAGGGAA GGTTATCTT GAGGGAGTCG	840
ATATTACGGA CAAGAAGAAT GACCTGTTG CCATGCGTGA GAAGATGGC ATGGTTTTTC	900
AACAATTCAA TCTCTTCCT AATATGACTG TGATGGAAA TATCACCTTG TCCCCTATCA	960
AGACCAAAGG TGACAGTAAG GCCGTTGCAG AGAAAAGAGC TCAGGAACCTT TTGGAAAAAG	1020
TTGGTTGCC AGATAAGGCA GACGCTTATC CACAGAGTTT GTCAGGTGGC CAGCAACAGC	1080
GGATTGCCAT CGCGCGTGGG TTGGCTATGG AACCAGATGT TTTGCTCTT GACGAGCCAA	1140

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CTTCAGCCCT AGATCCTGAG ATGGTTGGAG AAGTTCTGGC TGTTATGCAA GATCTAGCCA	1200
AGTCAGGAAT GACCATGGTT ATCGTAACAC ATGAGATGGG ATTTGCCGT GAGGTGGCAG	1260
ATCGTGTCACT CTTTATGGCA GACGGTGTGG TTGTTGAAGA CGGAACACCT GAGCAGATT	1320
TTGAACAAAC CCAAGGACAA AGGACTAAAG ACTTCTTGAG TAAGGTTTA TAAGTTAGCT	1380
TTGTTTAGCT ATTTGTAGCC AGCTTTAAC AC TAAAGAGA AGATTAGTGA AAAGCTAAC	1440
CAGAGCTTT TCTTATAGTT TAAAGCTATA GGATTGCCTA GGAAAGAAAGT GTTAGAGCTA	1500
CATTGTATTT TTTGGTATAA TTAAAGATAT TTGTAAGAAA AGAGAAGTGA TATGACACAG	1560
ATTATTGATG GGAAAGCTTT AGCGGCCAAA TTGCAGGGGC AGTTGGCTGA AAAGACTGCA	1620
AAATTAAAGG AAGAACACAGG TCTAGTGCCT GGTTGGTAG TGATTTGGT TGGGGACAAT	1680
CCAGCCAGCC AAGTCTACGT TCGCAACAAG GAGAGGTCAG CCCTTGCGGC TGGTTCCGT	1740
AGCGAAGTAG TACGGGTCC AGAGACCATT ACTCAAGAGG AATTGTTAGA CCTGATTGCT	1800
AAATACAATC AGGATCCAGC TTGGCATGGG ATTTGGTTTC AGTTGCCATT ACCAAAACAC	1860
ATTGATGAAG AGGCGGTTCT ATTGGCTATT GACCCAGAAA AGGATGTGGA TGGTTCCAT	1920
CCTCTAAACA TGGGGCGTCT TTGGTCTGGT CATCCAGTC TGATTCCTTC GACACCGCA	1980
GGAATTATGG AAATGTTCCA TGAATATGGG ATTGACTTGG AAGGTAAAAA TGCAGTCGTC	2040
ATCGGTGAT CCAATATTGT CGGAAAACCT ATGGCCCAGC TTCTTTGGC AAAGAATGCA	2100
ACAGTAACCT TGACTCACTC ACGTACTCAT AATCTTCCA AGGTGGCTGC AAAAGCAGAT	2160
ATTCTGGTTG TTGCAATCGG TCGTGCCAAG TTTGTGACTG CTGACTTTGT CAAACCAGGT	2220
GCGGTAGTCA TTGACGTTGG GATGAACCGC GATGAAAATG GTAAGCTCTG TGGGGATGTT	2280
GATTATGAGG CGGTTCCCC ACTTGCTAGC CACATTACGC CAGTCCTGG AGGTGTCGGT	2340
CCTATGACCA TTACTATGCT GATGGAGCAA ACCTATCAGG CAGCACTTAG GACATTGGAT	2400
AGAAAATAAG ATAAAAATTT TCTGAGGAAA GTGTATTTTC TATAGCTATA TCTAAAATGA	2460
TAGAAATGAA TATTAATTT TAGAAATAAG TTTATAAAAG GAGGTTTGCG CCTCCTTTT	2520
GTTGTATAAT GGAGTGAGGT GATTAGATGA TTTTAAAAAT TTATAATGGG GAATATAGTT	2580
TACAATGGGA TGGAATATAC TACCTAGCAC TAATTGATTA TCCAAATATT CAAGAGTGGG	2640
AATTAGAAA AATTGCTAAA TTTATAGCTT ACGAAAAACT TCATAAACGT CAAACAAAGTA	2700
TTGAGTGTGC TGATTCTGT TTAAAAAAAG AAATTTAGA TTACATCTGT CAGCATCCCT	2760
TTCTGCCACC ATTTACTCCT ACAGATAAAA GAGTAGCCTC GACTTATGAC CTACATAAGA	2820
GGTTAGTGAC TTCAGACTAC TGTAGTCATA CTACGACTAT AGATGCAGCG ATTTCTATTT	2880

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TTAAAACCTGG	TCGTCTTTA	TCTGCTGTGA	AAGCCTTTGG	GCGAGATGCT	GAGGAGTTGG	2940
TTTGGGATAG	TCGAAATGCT	GCATCTGATC	CGATAGATTA	TTTGACTAT	GTCATGTTAG	3000
GGTGGTCAAA	TACAAGTTCT	GGTTATCGAT	TGGCGATGGA	GCGTTATTA	GGTCGAGCTC	3060
CTTCAGAGAA	AGAATTACAA	GACAAGTTA	TCCCTGGAGT	AAGTTTCAT	TTTATCTATA	3120
CAGATTTGAT	TAAAGTCCT	GGTTATATTT	TTGATGGTTA	CCATGCTGTA	AAAATTAAGG	3180
ACATGCTTAA	TTTATTAAGT	GAGTTGTATA	TTTGCATTAT	TCCAACTCAT	AATAAGAGCC	3240
AATTTGAAAA	TATTATTCCA	ACCAAAATAC	AAGATAGGGT	GTATTATCTT	GACTATGCTG	3300
GAGAAGACTT	AGAAGAGTGG	ACTAAGAAAG	TCTATCAAGT	TCTTTAAAAA	CAATCAGATA	3360
AAGGATAGTT	GAGGAAAAAA	CGATGAAAGT	GATTGATCAA	ACCTTACTAG	AAAAAGTCAT	3420
TATTGAACGT	TCTTGTACAA	GTCATAAAGG	AGACTACGGT	CGTCTGCTGT	TGCTTGGTGG	3480
GACTTATCCT	TATGGTGGTG	CCATCATCAT	GGCTGTTTA	GCAGCTGTAA	AAAGCGGTGC	3540
AGGATTGGTA	ACCGTTGGAA	CGGACAGGGAA	AAATATCCCT	GCTCTACACA	GCCATTTGCC	3600
TGAGGCTATG	GCCTTTCTC	TGCAAGATCA	GTAATTGTTA	CAAGAGCAAT	TGGAGAAGGC	3660
AGAAGTTGTC	TTGCTGGGC	CTGCTTACG	AGACGATACG	TTTGGAGAAA	ATCTTGTAAA	3720
ACAGGTCTTT	GCTAGCTTAA	AAAAGAATCA	GATTTTGATT	GTAGATGGAG	GGGCCTTAAC	3780
CATCCTTGCT	AGGACAAGTT	TGTTGTTCC	ATCTAACAG	CTTATCTTAA	CTCCCCACCA	3840
AAAAGAATGG	GAAAAACTGT	CTGGTATTGC	TATTGAAAAG	CAAAACGAAG	GTACAACATC	3900
TAGTGCCCTG	ACTTCTTCC	CTCAAGGAAC	AATTTGGTA	GAGAAAGGTC	CAGCTACTCG	3960
TATTTGGCAA	GTTGGCCAGT	CTGATTATTA	CCAGTTAAAG	GTTGGCGGTC	CCTATCAGGC	4020
GACTGGTGGT	ATGGGTGATA	CACTGGCTGG	AATGATTGCA	GGATTTGCAG	GCCAATTCG	4080
ACAGGCCAGT	CTCTACGAAC	GTGTGGCAGT	AGCAACCCAT	CTTCATTCA	CCATAGCCCA	4140
AGAACTATCT	CAAGAAAATT	ATGTTGCTT	GCCGACGGAA	ATTAGTAATT	GTCTTCTAA	4200
AGTAATGAAA	AGATATGTCT	AAAATAGTTA	GACAAAAAT	GTTGATAATT	TGTATCATTA	4260
TTCTTAATTC	ACAAAAAACG	AACGTTTAGT	ATTCTTCTTG	CTAAGAAACT	AAATTGTTTC	4320
GTTTTTTTAC	TCTTGTAAAT	CTATTTTGT	TAGAGTTGAT	TTGGTTTACA	TCCGTACTTA	4380
AATTGATTG	TTAGAGCTCT	ACTTTATTA	AAAAAATTCA	ATTCAGGAA	TAAATAAGCA	4440
GTATTCTAAA	GGTACTTTA	GATGAAATAA	AAGCCTTAC	ATGGTATAAT	AGAGGTAGCT	4500
CTTTAATGGA	GGTGTGAG	TGGAAAATCT	GAAGAAAATG	GCAGGTATCA	CGGCTGCTGA	4560
ATTTATCAAG	GATGGGATGG	TTGTAGGGCT	AGGAACAGGT	TCTACTGCCT	ATTATTTGTT	4620
CGAAGAAATC	GGTCGTCGAA	TCAAGGAAGA	AGGCTTGCAG	ATTACAGCTG	TGACGACTTC	4680

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TAGTGTGACC	AGTAAACAGG	CTGAAGGGCT	CAATATCCCG	CTCAAGTCTA	TTGACCAAGT	4740		
AGACTTTGTC	GATGTGACAG	TCGACGGGGC	GGATGAAGTG	GATAAGTCAGT	TTAATGGAAT	4800		
CAAAGGCGGT	GGTGGTGC	CCC TTCTCATGGA	AAAGGTGGTC	GCAACACCAT	CAAAAGAATA	4860		
CATTTGGGTG	GTGGATGAAA	GCAAGCTGGT	CGAAAAACTA	GGTGCTTTA	AATTGCCAGT	4920		
AGAACTGGTT	CAGTATGGT	CAGAGCAGGT	CTTTCGTCT	TTTGAACGAG	CTGGCTACAA	4980		
ACCAAGTTTC	CGTGAAAAG	ACGGCCAACG	TTTGTGACC	GATATGCAGA	ATTTTATCAT	5040		
TGACCTCGCC	TTGGATGTCA	TTGAAAATCC	AATTGCTTT	GGACAAGAAT	TGGACCATGT	5100		
CGTTGGTGT	GTGGAGCATG	GT	TTTATTCAA	CCAATGGTG	GATAAGTAA	TCGTTGCTGG	5160	
ACGAGATGGA	GTTCAGATT	CAACTTCAAA	AAAAGGAAAA	TAGAAGGGGG	CATAAGATGT	5220		
CTAAATTAA	TCGTATT	CAT	TTGGTGGTAC	TGGATTCTGT	AGGAATCGGT	GCAGCACCAG	5280	
ATGCTAATAA	CTTTGTC	CAAT	GCAGGGTTC	CAGATGGAGC	TTCTGACACA	CTGGGACACA	5340	
TTTCAAAAC	AGTTGGTTG	AATGCCC	AA	ACATGGCTAA	AATAGGTCTT	GGAAATATT	5400	
CTCGTAAAC	TCCTCTTAAG	ACTGTAGCAG	CTGAAAGCAA	TCCA	ACTGG	TATGCAACAA	5460	
AATTAGAGGA	AGTATCTCTT	GGTAAGGATA	CTATGACTGG	ACACTGGAA	ATCATGGAC	5520		
TCAACATTAC	TGAGCCTTTC	GATACTT	C	GAACCGGATT	CCCAGAAGAA	ATCCTGACAA	5580	
AAATCGAAGA	ATTCTCAGGA	CGCAAGGTTA	TTCTGAA	GC	AAACAAACCT	TATTCA	5640	
CGGCTGTTAT	CTATGATTT	GGACCACGTC	AGATGGAAAC	TGGAGAGTTG	ATTATCTATA	5700		
CTTCAGCTGA	CCCTGTTTG	CAGATTGCTG	CCCACGAAGA	CATTATT	CC	TTGGATGAAT	5760	
TGTACCGTAT	CTGTGAATAC	GCTCGTTCGA	TTACCC	CTGA	GCGCCTG	CC	TTGCTGGTC	5820
GCATCATTGC	TCGCC	TTAT	GTAGGTGAAC	CAGGTA	ACTCGTACG	GCAAA	CCGTC	5880
GTGACTTGGC	TGTATCTCCA	TTTTCCC	AA	CTGTTTGGA	TAAATTGAAT	GAGG	GCTGGTA	5940
TCGATACTTA	TGCTGTGGGT	AAAATCAACG	ATATCTTAA	CGGTG	C	TG	ATCAACCATG	6000
ACATGGGTCA	CAACAAGTCA	AATAGTCATG	GAATTGATAC	ACTATTGAA	ACTATGGAC	6060		
TTGCTGAGTT	TGAAAAGGA	TTCTCATTCA	CAAACCTAGT	TGACTTTGAT	GCC	CTTACG	6120	
GCCATCGTCG	TAATGCTCAC	GGTTACCGTG	ATTGCTTGCA	TGAGTTTGAT	GAAC	GCTTAC	6180	
CTGAAATTAT	CGCAGCTATG	AGAGAGAATG	ACCTCTCTT	GATTACTGCG	GAC	CATGGAA	6240	
ATGACCCAAC	GTATGCCAGGA	ACGGATCACA	CTCGGGAAATA	TATTCCATTG	TTGGC	CCTATA	6300	
GCCCTGCCTT	TAAAGGAAAT	GGTCTCATT	CAGTAGGACA	TTTTGCAGAT	ATTTCAGCGA	6360		
CTGTTGCCGA	TAAC	TTGGGT	GTGGAAACTG	CTATGATTGG	GGAAAGTTTC	TTAGATAAAT	6420	

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TGGTATAAGA	TGACCGCGCTA	TGCCTTGCTG	G TGAGAGGTA	TCAATGTTGG	TGGTAAGAAT	6480
AAGGTCGTCA	TGGCGGAGCT	TCGTCAAGAA	TTGACAAACT	TGGGACTGGA	AAAGGTTGAG	6540
AGCTACATCA	ATAGTGGCAA	TATTTTCTTT	ACTTCGATAG	ATTCCAAAGC	CCAATTGGTT	6600
GAAAAGCTAG	AGACTTTCTT	TGCAGTCCAT	TATCCATTAA	TTCAGAGCTT	TTCTTTACTG	6660
AGTCTAGAGG	ACTTGAGGC	GGAACTTGAA	AATCTACCAG	CTTGGTGGAG	CAGAGACTTG	6720
GCACGAAAAG	ATTTTCTCTT	TTACACTGAG	GGTTTGGATG	TGGACCAAGT	CATCGCGACA	6780
GTTGAAAGTT	TAGAGCTGAA	AGATGAAGTG	CTTTATTTG	GAAAAC TTGG	GATTTCTGG	6840
GGGAAATTTT	CTGAAGAAC	CTATTCTAA	ACTGCCTATC	ATAAGTACTT	GCTGAAGGTG	6900
CCTTTCTACC	GCCACATTAC	TATTCGTAAT	GCTAAAACCT	TTGACAAAT	TGGTCAAATG	6960
CTAAAAAAAT	AATAAAGGAG	ACACACAATG	ACATTTTTAA	ACAAAATCCA	TGAAAC TGCT	7020
ACTTTCCCTGA	AAGAAAAGGG	AATTGCAGCC	CCTGAGTTCG	GTCTAACCT	TGGATCAGGA	7080
CTTGGAGAAT	TGGCAGAAGA	AATCGAAAAT	CCAGTTGTAG	TAGACTATGC	TGAGATTCCA	7140
AACTGGGGCC	GTTCAACAGT	AGTCGGTCAT	GCTGGTAAAT	TGGTATATGG	TGAAC TGCGA	7200
GGTCGCAAGG	TCTTGGCTCT	TCAAGGGCGT	TTCCATTCT	ATGAAGGGAA	TCCTCTGGAA	7260
GTGGTGACTT	TCCCAGTTCG	TGTGATGAAA	GTTCTTGGAT	G TGAAGGTGT	TATTGTAACC	7320
AATGCAGCTG	CGGGTATCGG	ATTTGGTCCT	GGTACCTTGA	TGGCTATCTC	AGACCATATC	7380
AACATGACGG	GGCAAAATCC	ATTGATGGGT	GAAAAC TTGG	ATGACTTTG	CCCACGTTTC	7440
CCAGATATGT	CTAGGGCCTA	CACACCAGAA	TACCGTGCCA	CTGCCCATGA	AGTGGCTAAA	7500
AAACTTAATA	TCAAGCTTGA	TGAAGGTGTC	TATATCGGAG	TTACTGGTCC	GACTTATGAA	7560
ACACCAGCAG	AAATT CGTTC	CTATAAGACA	CTGGGAGCAG	ATGCAGTTGG	TATGTC TACG	7620
GTTCCCTGAAG	TTATCGTGGC	AGCCC ACTCT	GGCTTGAAAG	TTCTGGGAAT	TTCATGTATC	7680
ACTAACTTTG	CGGCCGGTTT	CCAAGAAGAA	CTCAATCACG	AAGAAGTTGT	AGAAGTGACT	7740
GAACGTGTTA	AAGGTGATTT	CAAAGGCTTG	CTTAAAGCGA	TTCTTGCTGA	ATTGTAAGAA	7800
AAAAGATTAA	AAAGGGGGAG	TGCCTCTGTT	TTTCAGGAT	TGACTGCTA	TCCGGATTAA	7860
AGAAGAAACA	GAGGAATACT	ATGAGCTTCT	TCCTGCTCTT	ATAACTGAAA	GAAGCGGAAG	7920
AATAGGTATG	TCTGATCTGA	TAGCCAGCAT	TGTGAAAGAC	AAGATTCTAG	GATACTAGCA	7980
TTAGCTTCCCT	AGCCAAGCAG	ACTAGTATGA	TAAGGAGAGA	TGAGAATGAA	TTGACTTTCT	8040
GAATTTCTCA	GTCTTATCAT	ATATAGCACA	ATGAGATTT	GCTTGAGTCT	GCTTGAAAT	8100
AAACGAAAAG	AAAGATAAGA	AATAATGAAA	ATTGGTCAAC	GAATTATGCG	CTTTGGCATA	8160
AAAAATTAAAG	TATCGGAGTT	GTATCTGTTG	TAGTCGGCTT	TGATTTCTAG	CTCCAGCTGG	8220

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AATTCAGCC	AATGAAGTAA	AGCAAGATGT	AACATCTGAA	GTGGTAATAG	GTGTGCTAGA	8280
TTCTAAGGAG	GAATTGAAAG	AGTCAGAAAA	TGATGCTCCA	AAACTAGAAA	CTCCTCTTAG	8340
AGAGGAGCCA	AGACTAGCTC	CTCAAACGCT	TCCGGAAGCA	AGTGAAGTTC	TTGAAAACAA	8400
AAGGGAAGAG	TCAAAAGTAG	AGATAACATA	ACCAGCTCAA	GCGGATGATA	TCCGCAAGGT	8460
TGTTGGGAA	TTAGCCAAGG	ATATAAGTAT	TACTAAGTTG	TATATGACAG	GTCATTCTCT	8520
TGGATGTTAC	CTAGCTCAGA	TTGCAGCGGT	TGAAGCTTAC	CAAAAATATC	CTGATTTTA	8580
TAACCATGTA	TTGAGGAAAG	TGACAACTTT	CAGTGCTCCT	AAAGTGATTA	CTTCCAGAAC	8640
TGTTTGGAAAT	GCTAAGAATG	GTTCCTGGGA	TGTTGGTTTG	GAAAGTCGTA	AATTAGCTGT	8700
TAGTGGAAA	ATTAAGCATT	ATGTGGTTGA	TAATGACAAT	GTTGTGACTC	CCTTGATTCA	8760
TAATAATCGT	GATATTGTTA	CATTACAGG	TAATTCACGC	TTTAAACACC	GTTCTCGTGG	8820
CTATTTGAA	AGTCCAATGA	ATGATATTCC	TAACTTTAAT	ATTGGTAAAC	AAGCTACCTT	8880
GGATAAACAT	GGTTATCGTG	ATCCGAAATT	GGATAAAAGTG	CGATTCTTTA	AGAAACAGGC	8940
TCTGCCTCGA	TCTTCTAGTC	AACCAAGCGC	TGAACCAATG	GAAAATATTG	CCTCAGGAAA	9000
ACAGGTTACT	CAAAGTTCGA	CAGCTTCGG	AGGAGATGCT	AGAAGAGCTG	TGGATGGCAA	9060
AGTCGATGGT	AACTATGGTC	ACAATTCTGT	CACTCATACA	AACTTCCAAT	CTAACGCTTG	9120
GTGCCAAGTA	GATTGGCTA	AAGAAGAAAC	CATTGCCAA	ATCAATATT	ACAACCGAAC	9180
AGACACTGCC	CAGGATAGAT	TGGCAAACCTT	TGATGTCATT	CTTTTAGACA	GTTCTGGTAA	9240
AGAAATTGAG	TGAAAACGTA	TAACATCTCC	TAAAGATGTG	TCAGCACAAA	TTACGATTAA	9300
CCATAAAAAA	CCGCGCTATG	TTCGGATTGA	GCTAGAAGGC	TATAATGCC	TCAGTCTTGC	9360
AGAAGTTGAA	GTTCCTGCT	TTATAGCTAC	GAATGCTGAA	ACGGCGACAC	AAGTTCTAA	9420
GCCAGTTCAA	CCAATCAGTC	AGACTCCTGT	GAAGGATAAA	ACATTGACAA	TTCAACACAG	9480
TGGAGCTTAC	ATTGCCCGCT	ACTCCATAAC	TTGGGAAGAA	GTTCCAGTAG	ATAAAGATGG	9540
AAACCAAGTT	GTTCGTAGTC	ATTCTTGGGA	AGGAAGCGGT	CGCAACCAGA	CTGCAGGTT	9600
TGTCCTCAAC	CTCCCAATCA	AAGAAAATAT	GAGAAATCTG	CGAGTTAAGA	TTGAGAAAAAA	9660
GACGGGCCTA	CTATGGAATA	GATGGCAAAC	AATCTATGAA	AACAGACCAA	TTTTAGCTCA	9720
ACCCCACCGT	AAAATTACCC	ATTGGGGTAC	GACATTGAAT	TCCAAGGTGA	GTGACGATGA	9780
TGTCTTGTAA	TCTGATGGTA	GAATGACAGT	TAGTTGTCT	AGTTTATAAG	AAAGTACTAC	9840
CTGAGCTTGA	ATAGGACTCA	GGTAGCTCTC	TATGAAAGAA	CAAAATTAAT	ACTCAATGAA	9900
AATCAAAGAG	CAAACTAAGA	AACTAGCCGC	AGGTTGCTCA	AAGCACTGCT	TTGAGGTTGT	9960

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AGATAAGACT GACGAAGTCA GTCACATATA TAATCCAAGG CGACGTTGAC GTGGTTTGAA	10020
GAGATTTCG AAGAGTATAA ACAGAAAGGT AGAGCGCGTG TTCTAATTG AACACGAGTA	10080
GAAAACTTT CTAAAACAA AAACGAAAGG ATGGGTAAAC TGTATTCGCT GAACTGAATA	10140
CGGGCGACTC TCCTCTAAAT CAAAATTAAG AAAGGAATTG ACCCCCACCT AAAAGTAGTG	10200
GGAAAAAGAT AGTTGATCTA GCGAGCATCG CTCACTGCAC CCAACTCCTA TTTCCCTTC	10260
GCTTTTGTG GGGTTTGGTA TCTTCTCAA TATAAAATAT AAAATAAAGA AAGGTAGAGC	10320
GTGTGTTTG ATTTGAACAC GAGCGGAAAA CTCGGAAAAT AGATAATCTG ACTGAAAAAT	10380
CAGGATTTCT CGTCAGGTTC CTAATTTCA GTCGTTTCT TCTCGCTCT TGTATCATAA	10440
ATTATGTCTA TCCATATTGC TGCTCAGCAG GGTGAAATTG CTGATAAAAT TCTTCTTCCT	10500
GGGGATCCTC TTCGTGCTAA GTTTATTGCG GAGAATTTC TTGATGATGC TGTTTGTGTT	10560
AACGAAGTGC GTAACATGTT TGGTTACACT GGTACTTACA AGGGTCACTG TGTATCTGTC	10620
ATGGGAACTG GGATGGGAAT GCCATCTATT TCGATTTATG CGCGTGAGTT AATCGTAGAC	10680
TACGGTGTGA AGAAATTGAT TCGTGTGGGA ACTGCAGGTT CTTTGAATGA AGAGGTTCAT	10740
GTTCGTGAAT TAGTTTGGC GCAGGGGGCT GCAACCAACT CAAACATCGT TCGTAATGAC	10800
TGGCCACAGT ACGATTTCC ACAAAATTGCT AGCTTGATT TGCTTGATAA AGCCTACCAC	10860
ATCGCCAAAA AACTTGGTAT GACTACTCAC GTTGGGAACG TTTTGTAC TGTATGTCTT	10920
TACTCAAATT ACTTTGAAA GAATATCGAG CTTGGTAAAT GGGGAGTCG GGCTGTGGAA	10980
ATGGAAGCAG CAGCTCTTA CTATCTGCT GCCCAATACC ATGTTGATGC GCTAGCTATC	11040
ATGACCATCT CTGATAGCTT GGTCAATCCA GACGAAGACA CAACTGCAGA AGAACGTCAA	11100
AATACCTTCA CTGATATGAT GAAGGTTGGT TTGGAAACCT TGATTGCGAG ATAATTATAG	11160
CCAAAAAGGG GCTCTTGTC AACTGTAGTG GGTTGAAAAA AAGCTAAGCT TGAGAAAGGA	11220
CAAATTCGTTT CCTTTCTTT TTGATATTCA GGGCGATAAA AATCCGTTT TTGAAGTTT	11280
CAAAGTTCCG AAAACCAAAG GCATGCGCT TGATAAGTTT GATGAGATTA TTGGTCGCTT	11340
CCAGTTGGC ATTAGAATAG TGTAGTTGAA GGGCGTTGAC GATTTCTCT TTGTTCTTA	11400
GAAAGGTTTT AAAGACAGTC TGAAAAAGAG GATGAACCTG CTTCAGATTG TCCTCAATGA	11460
GTCCGAAAAA TTTCTCAGGG TCTTGTCT GAAAGTGAAA AAGTAAGAGT TGATAGATCT	11520
GATAGTGGTG TTTCAAGTCT TCTGAATAGC TTAAATCTT GTCAAGAATT TCTTATTTG	11580
TTAAGTGCAT GCGAAAAGTA GGGCGATAAA AACGTTTATC GCTsArTTA CGACTATCCT	11640
GTTGGATGAG TTTCCAGTAA CGCTGTAGAG CCTTGTATTG ATGAGATTT CGTTCAAAC	11700
GATTCTATAAT TTGAACACGA AAACGACTCA TGGCACGGCT GAGATGTTGG ATAATATGGA	11760

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AACGATCTAG AACGATTAA GCACACGGAA AAAGCTGTT AGCCAAGTCA TAGTAAGGAC	11820
TAAACATATC CATCGTAATG ATTTCACTT GACAACGAAC GGCTCTATCG TAGCGAAGAA	11880
AGTGATTCG GATGACAGCT TGTGTTCTGC CTTCAAGAAC AGTGATAATA TTAAGATTAT	11940
CAAATCTTG CGCAATGAAA CTCATCTTC CCTTAGTGAA GGCATACTCA TCCCAAGACA	12000
TAATCTTGG AAGCCGAGAA AAATCATGCT CAAAGTGAAA GTCATTGAGC TTGCGAATGA	12060
CAGTTGAAGT TGAAATGCC AGCTGATGGG CAATATCAGT CATAGAAATT TTTCAATTA	12120
ACTTTGAGC AATTTTTGG TTGATGATAC GAGGGATTTG GTGATTTTC TTTACCAGGG	12180
GAGTCTCAGC AACCATCATT TTTGAAsAGT GATAGCACTT GAAACGGCGT TTTCTAAGGA	12240
GAATTCTAGA AGGCATACCA GTTGTTCGA GGTAAGGGAT CTTAGACGGT TTTGAAAGT	12300
CATrTTTCTT CATTAGACTT CCACAATCAG GGCAAGATGG AGCCTCATAA TCCAGCTTAG	12360
CGATAATTTC TTTGTGGTA TCCATATTGA TGATATCTAG AATCTTGATG TTTGGTCTT	12420
TAATATCGAG CAGTTTGTG ATAAAATGTA ATTGTTCCAT ATGATTCTTT CTAATGAGTT	12480
GTTTGTGCGC TTTTCATTAT AGGTCAATATG GGACTTTTT TCTACACAAA AATAGGCTCC	12540
ATAATATCTA TAGTGGATT ACCCACTACA AATATTATAG AGCCAAAAA GGAAGCCCTT	12600
TATGAATTGT AGGACTTCCT TTTCTTATCC AGAAATTGAT CTAGCTCT CTGATTTCGA	12660
AGAATAGTGA CTTTATGTGA ATATTCTTGG CAAAGTTTT GGTAATTTC TTTTTGAGTT	12720
TTGCGGACGC CCATCCAAA GAATCCATCT GATAAACTCC CACTCAAAGC GTTCAGGGCA	12780
ATCTACCGCC ATACTTCTC TGACTTTCC ACGGTATTTA AGATAACGCT TAAAGGCTCT	12840
AAAGAGACAG GTCAATGCC AAAAATTGAG AAAGATGATT TGGTCAGCTT CTTGCATTG	12900
TTCTTGGTAG TAGCACCAAG AATAATTACC ATCGATGACC CAAGTTTAT GCTTGGTGAG	12960
AAAGTTTTT ATCTCGTTA ACATCCATTC GCAGTCACTG TCTTGCCAAAC CAGGTTGAAA	13020
TTGGAGTGTG TCCATGTGCA GTTTGGAAT GGAGTAGTAG TTAGATAACT TTTCTGCTAT	13080
AGTTGACTTA CCAGAACAG AATATCCGAT AATTGCGATT TTCATTTCT ACCTTTCCCT	13140
ATTTGGAGAC AAAAACACAG CCTCTATGGA CTGTTCTTA TTTAACAGT TTAGCTGAAA	13200
GACGAGCTTT ATCGCGCTT GCTTGTGTT TGTGAATCAA ACCTTTAGTT TCTGCTTTAT	13260
CGATAGCTGA GCTAGCAGCA CGGAAAAGTT CTTCAGATGG GTTGCCTTCG AAAGCTTTA	13320
TAGCAGTACG CATAGCTGAT TTTTGAGCTG AGTTCTTTTC GATTCGTCTA ACGTTCAATT	13380
CAGCGCGTTT GATAGCTGAT TTAATGTTG CCAATGGTCT TACCTCCATA TTTACTAACT	13440

(2) INFORMATION FOR SEQ ID NO: 129:

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(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8512 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 129:

CCTTTTTTCA	AAAACTAGAT	ACTAGTCTAT	CAAAAGTAGG	AAAGGGTTTC	AAGAAAATTG	60
ATTGGAAATT	TTTGAAAT	CATAGAACTA	TTAGCTAATC	CCTAGTATTG	AAAAGACTGG	120
ATAGCTTCTT	TCAGGTCA	TC	TTGTAAACTA	TTCTCTGGT	CAAGTTGGAC	180
ACCAGACAGG	ATCTAAAGTT	GGAAAATTG	AAAAAATCCT	CCCTTCTTC	TATCGGAAAA	240
TCAACAGTTT	TTATCCAAGA	AGCTACTTGT	TCTTGCTCCA	ACTTCCCCTG	TAAAATAGGT	300
TCATAGATCA	CTCTGCTAA	ACGCCAATCC	TCATCATCTG	TAAAGCGAAC	CGACATTCTT	360
TTAAATAGTT	GGCCAAGTAT	ATCAAATACT	TCATGAAC	TGTTTTAGG	AAAGTCTGGA	420
TGACAAACCA	CCTCTGTCAG	TAATCGGCT	CCATGTGCAA	AAGCGTGAAC	CCAACCATAC	480
TGACTTGAGA	AACCCCTTGT	ATCCTTTCT	TTTGAAAGAT	AGTGCAAGCC	TTGATTAAA	540
AGGACATTAC	GAATTCTGG	AGAAGGATTT	CCCAAATGAT	CAAACAACCA	CTGGATTCT	600
TCCTGGTTAT	AATTGGTTT	TTCTTCTGCT	ATTTTCTTA	GTAAATCTG	ATACATGGTC	660
AATACCTCTA	CATTTCTAGC	AACTGTTCAA	AAAGGCAGTC	TTAAATGACT	CAATATTGAA	720
TTCTCAATTA	AATACAATCT	GATATAAAAT	GACGTAAATA	ACTATCAATA	CCAGTTCTAC	780
AGTAAGTTCA	AATTAAACAT	CACGACCTTC	AACGACATT	TTGAAAATAG	CTACAAC	840
GACAAATAGA	ATGACGCTTA	ACAAGCCC	AAACATCATT	CTAAAAAATT	TTTCTATTCC	900
CCTACTCTCC	CAACTCAGCA	CTATAGGAGA	TAATCTGGTC	AACTGTGTCA	GACAAGAATT	960
GGATGGTATC	ACGGAGTGGT	TTGCTGTTG	AAATATCAGC	ACCGATAATC	ATGGCTGACT	1020
CAAGTGGTGT	CTTGCTACCA	CCTGATTGAG	GGAGATTGAG	CCAGTCTTCA	GCTCCAGTTT	1080
CAGAATGTTT	TAGATGAAGG	TAACCAGCAG	TCGAGATAAC	TAGTCCTGCT	GACTAAGTGT	1140
AACTATACAA	GCCCCATATAG	TAGTGAGCTT	GGCGCATCCA	AGTCAGAGTT	GCATCATCGT	1200
CAATTTCAT	AGCATCTCCC	CAGAAATCCG	TCAAAACTTC	CTTCATAATG	CTGTTGAGCT	1260
TGCTTGCTCC	AAAGGTCTCC	CCTTCTTCAA	TCAATGTATA	CACCTTACGC	TGGAAGGCGG	1320
CTTCCAAGAG	GTGGGTGATA	AAGTTATGGA	AGTAGGTGTC	TGTCAAGCGA	TGAGCCAGAG	1380
CGAAGCGTTT	TTGACGTGGG	TCATTAGACT	GGTTCTCCAA	GTAATCACTG	AGTAGCAATT	1440
CATTGAAGGT	TGACGGTGCT	TCAACATAGT	AGGTGACAT	ATGGGCATTG	AAGTAAC	1500

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GATGATTGTC	TGAAAAGATG	AATTGACCAG	AATGCCGAT	TTCATGAATC	AAGGTATAGA	1560
CATCGCTCAA	ACGGCCTGTC	CAGCTCATGA	GTACATAAGG	GTGTACGCGA	TATGGTCCG	1620
CCGCATAACC	ACCGGAATCC	TTGCCACTGT	TAGCAGCAA	GTCCACCCAG	CGCTCTCTT	1680
GGTAACGAGC	AACTTCCTGA	CAATATTCTT	GCCCCAAAGG	TTCTACCGAC	TTCATGACCA	1740
AATCATAGGC	ATCGTCAATA	GTCACTTCAG	GATTCAGGGC	GCTGTCCAAG	TCCAATTCC	1800
AGTCTGCAA	GGTCATCTT	TCAAGACCAT	TTACCTTGTC	AACATGCTG	AGGTATCTCT	1860
GAGCGACTGG	TGCAAAGTCC	TTCATGATGA	GGTCAATCTG	GCGGTCAAAC	ATGACACGGT	1920
CCACCTCTTG	TTCAGCTAGA	AGATAGTCAA	AGACAGAGTC	GTATCCCTTC	ATATCAGCCA	1980
AGAGTTTTTC	AGACTTGACC	TGAGCCAGAT	AGGCTGCTGC	AGCCGTATTT	TGGTGCTTAC	2040
GAAGTCCCTC	TGAGAAGGAA	CGGAAGGATT	TCTCACGAAC	CTCAGCATCC	TCATGGTTTT	2100
GGTAGAAATT	CTCATAGGTC	ACAAAGCTGT	TTTGTTAGGT	CTTGCCATGG	GCTTCAAAGT	2160
CAGCCATTTC	AAAATCCCCA	GCTCGCATCT	TAGTATAAAAT	GTCCTGCGGA	CTGTAGAAAA	2220
CTTCACCGAG	ATTTGTCAAG	GCCTTCTCCA	CATCTGCC	TAAGTAGTGG	GCTTTTTGGA	2280
TTTTAGCCTG	ACGAATGGCA	GCTGTTAAAT	GTGGCAATT	ACCCAAACGG	TCCAAGACTT	2340
CCTCATCTGC	TGCCACCAAG	GCATCGTCAA	AGAAGGTCAA	GGCTACGCTG	GCATCTGTT	2400
CAAATTCCAT	CCCAGCTTGG	GCAATATTGG	CAAATTGTC	ATTGCTATAG	TCCGTCGTCT	2460
GAGGCATAAA	ACCATAGTTG	CCAATATGGC	TCATCTGAAT	GTAGATCTGT	TCCAATTCCG	2520
CAAAGGCCTT	CTCGAAATCC	TCAAAAGCTGT	GAAGATTGCC	CTTGTAAATCA	CGGCTAAACT	2580
GGTTGATGTC	TTCGCGAGCT	TTCTCGATTG	CACGCAAGAA	ATCCTCACGG	TCTGGTATA	2640
GGGCTGTTAA	GTCCCAGAGT	TCCTTCTCTG	GAAATTCTGA	ACGGTGT	TGTTCCATT	2700
TCTTCCTCTT	ATTTCTCTAA	TTCTACTAAA	ACACTAAGGG	CTGATAAAC	GTAAAGCGGT	2760
GCTGTTCTG	CTCGAAATAT	ACGAGGACCT	AGGCCTGCCA	AAACGGCTCC	TTTAGCTTCA	2820
AAACTTTCGA	TTTCTGCAGG	TGAGAGACCG	CCTTCTGGAC	CAAAGATAAA	GAGCAGTTG	2880
GCTCCTGTT	CAAGACCAGT	GACTGCTTGC	AGAAGCGCAG	CGGCTTCTCC	TTCTTAGCT	2940
GATTCTTCAT	AGGCTACTAT	GATAGAGTCA	AACTGGTCCA	GCTGAGCTAG	AAAATCTGCT	3000
TTTTTCTCGA	AAAGTTTAAT	ACTTGGTACA	ATATTACGCT	TGCTTGTCTC	GGCTGCTCCA	3060
AGGGCAATT	TTTCTAGTTT	TTCAACTTTT	TTACCCAATT	TCTTGCCATC	CCACTTGGCA	3120
ACTGACCAGT	CTGCAGGAAA	GGCCCAGATT	TGGCTAGCCC	CCAGTTCGGT	TACTTTTGA	3180
GCGATGAACT	CCAGCTTGTGTC	TCCCTGGGA	AATCCAGATG	CGATGGTCAC	TTGGACTGGT	3240

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AGTCCACAT	TGTCATTAA	TTCTGGACC	AACTCAAAC	GACGATTTTC	CATATCCAGC	3300
ACGCGCGCCA	AGCGCTTGAT	GCCATCATCA	AAGACTAAGG	TAACCTCATC	CTCTCTTTC	3360
AAGCGATAA	CCTGAAACAT	ATGCTTACTG	GTTCCTTGT	CCTCGATAGT	GACAGGAGAG	3420
ATAGCACTGC	CTTTTACAAA	ATACTGCTGC	ATGCTAGCCT	CCAATCACAC	CAGAGATATC	3480
CTTGGTTTTC	TTAAAGACAC	AGGTATTCCA	TTCCCCTTGA	ACCATGTGAG	TTTCGAGGAA	3540
AAATCCAGCT	GAETCAGCCG	ACTGGCGCAC	CATGTCCAAC	TTGTCTTGA	TAATGCCACT	3600
CATGATCAGG	TAGCCTTCAT	CCTTTACCAA	GCGATAAGCA	TCGTCTATTA	GATGAATGAG	3660
GATATCCGCC	AAGATATTAG	CCACAATCAC	ATCTGCCTCA	ATTTCCACAC	CCTTAAGCAA	3720
ATCTCCAGCC	GCTACATGGA	TATTTCCAT	GCCAGGGTTG	AGCTCAATAT	TTTCCTGAGC	3780
CACACGAACC	GCCACATCAT	CCAGGTCATA	GGCGAAAATT	TCTTTAGCCC	CCAGAACGGA	3840
GCTGGCAATA	GAGAGAACCC	CTGAACCACT	CCCCACATCT	AGCACCGTTT	CGCCACCACG	3900
AAGAACCTGT	TCCAAGGCAA	AAAGGCTCAT	CTTGGTAGTT	GGGTGGGTTTC	CAGTACCAAA	3960
AGCCATGCCA	GGATCCAGCT	TGATAATCAT	TTCCCCCGCA	GTCGCCTCAT	AGTCTGTCCA	4020
AGAGGGAACG	ATGGTCAAAT	CATGAGTGAT	ACGAGCAGGT	TCATAGTATT	TCTTCCAGTT	4080
GTCTGCCAG	TCTTCCTCAG	CCAAGGCAGT	CGTACCTATT	TTTAACTCTC	CCAAATCCAT	4140
AAAATCTGTC	AATTCTGCTA	GACGAGCCTG	CAAATCCGCC	TCAACCACTG	TCACATCCAC	4200
CGTGTCAAGG	TAGTAGGCTG	TCACTACGAT	TTCTTCTTGC	TGCTCCACCT	CTGGGAAAAT	4260
CTCTCCAAAG	CGGTCCACAT	TTCCACATA	GTCCATACTG	TCTTCGATTG	CGACTCCTTG	4320
CGCTCCCAGC	TCAATCAAGA	GATTGGAAAC	CAACTCCTCT	CCCTCACGCT	TCACTGTAAAC	4380
TTTTAACTCT	TGCCATGTTT	CCATTATTAA	TACCAAGCCC	GTAAAACACA	AAACCAAAAT	4440
AGGAAATTCT	CTGAAGACGC	TTGTGTCTAA	GAGAAGTTA	TCTTTTGCG	ACAGTGTAA	4500
GGCGGGTTTC	AGTTTAGAAA	TGTAACGTAA	CCATCCTTTC	TAATCACTTA	CTTTAAATA	4560
ATCTTTAAAT	CTCTCTTGCA	ACTGAGGCAC	AACTTGACTG	GAACTAAGAA	ATTCCCTAAC	4620
ATTCACTCAGC	TGATAGCCCT	GTCCTTCATC	TCCGAAGATG	ATATTGTCAA	ATTGTTCTTG	4680
TCTTAGCTGA	CCAACCATAA	AGACCGATTT	CTTGCCTTTA	AAAATTACGC	TAGGATAAAAT	4740
CTTGCTCCAA	AGCAGACAGT	CTTCATCTAA	ATGAATTCCC	AGTTCCTCAT	AAACTTCACG	4800
CCGAGCGCAT	TCAAAAGGGC	TTTCGCCCC	TTCACGGCCA	CCACCTGGCA	GTTCCCACAT	4860
ATTGGCCAG	GGAATACTTG	CCTTATCATC	GCGTAAGATA	GTCAAAAGCT	TATCCCCACA	4920
AAACAAAGCA	ATCTTGCAAC	CTGTGAAATC	AGAAATTCT	AGTTCCTCATCT	TCAGTTCCCTT	4980
CTAACATTTC	CTTTTCCAGC	TCGGCTAAC	AGTTTCATA	ATATCTTTC	TCATCCCTCA	5040

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ACATTCGACT ACTATCCATT TTCTGTCTAG CAATCTTGAG AGCCTTACGA GTTCGATCTA	5100
CATCTTTCTT CACCTTTAAC TGATACCAGG CTTGTATCAC TTGAAGATTG GACAGTTGA	5160
GAGACAGAAA CGATTTGACC TGTGAAATAC TAGCATATTG CTCCGCTTGC TCAAATCTC	5220
CTTCCAACAA GGCGATATGA AGCAGGGATA GTTGGGCAAC TGTCTGCATC ATCGGAGTAG	5280
TTGTCCCTCTC AAGTAATGCT TGAAACTGCT GTTAGCTAC TTCTTCCTTC CCTTCCAAAA	5340
TGGAAACTTC ACCTTGACATA CCTAATACAC CATCCGCAAA ACTCCCTCGT GCATCCTCAG	5400
GAAC TGCTTG AACAAAGTCT TTCAAATCAT ATTCTTGAGG AGCTAGCAAG GTCTGGCAG	5460
AATGTCTCAA TACCAAGGAG GCGTATTGAG TATTTTCAGG GTGTTGTAGT AATTCCAAA	5520
TTTTTGCTCC ATCGGTGATG TCGACTGGCA AAATGTTATT TAGGAAGAAA GATAAATTAA	5580
GAAAAATCCA AGTCCCTGCA AAATACCAGC TTCTTGCAA AAATCCAAAC AATATGCCA	5640
ATAATATCAA GCCGAGATGA ACCATCAAGC CTCTGAAAG CATCAGGATG ATTCTTGAT	5700
CGCTTTCATC CTCTTTAAA CCAATGTTATT GAGCACCAAC ATTTTCAGA ATGGCTGTT	5760
TACTAAGATG AAACCTGCCT GACTTTTGAG TCAAAATAAA ATGTCCTAAT CCAAAAGCCA	5820
CCAGCCGATA GCCTGTCAAG TAGCCACAAA AAGCATGACC CAGCTCATGA AGAATAAAGA	5880
TTAAATACAT GCTTAGAAGA GCGAAGGCAT AACCAAAAGT AAAGGCTAAA ACTGCGGAAT	5940
ACCCCAACTC TGCAAATGCG ATTGTTCCAC AAGCAAAAGC TAGCATAATA AAGACAACAG	6000
CTAGCACATA AACCAAATAA GTCCAATTT TCTTCATAAC ACCTCCAACC AACTCCTAGT	6060
ATCTTGGATA AGGATAAAAT TCTCCCTTTT CCAAGCCAAT TTTTCCTTCT TCAAAGACTT	6120
CTTGGTTCCA TTCCATGACA AATTCCCTCTG CTTCTGGTC TTCCAAAAAG TCCATGAGGA	6180
CATCTAGCCC AACCTCAGCA GTATTTAA GGAAAAGCGC AAAATAAGCT AAAAATTAC	6240
GGGAAATCC TTTTTAGGC AGGTAAGGAA TAACAGTCAA ATAGTCTTCC TCATTGACTG	6300
TTGACTTGGC AGGATTGTAG AAAAGGACCG CTTCCCTAAA AAGAATGTCA TCTGATGAAA	6360
CCTCTCCGTC TTCATCCACC ATCTCCACAC CGCAGCATTG TGCGCTTCCA ATAGAAAAC	6420
CACTTCTACC GCATGGTGC GTTGTCCCA GCTAATCTCA AAGTCAAAGG GAAAGTTCTT	6480
GTCCAACCTCT CCCTCTAAAA TATCTAAAAA TCCGTATGTT GCCATTTGT CCTCTTTCTA	6540
TGCGACTCTT TAATCGCCCC GATTGCTCGG AAATATGCTA AAATAGATAC TACCATCTTA	6600
CCACAAAATT ATTTTATGTC CTAATTATAC CATATTACCT CATTAAACC CTTGGTATCA	6660
GTGATTTCT TAAAAGTCTG ATTTCTTCAT TTCTCATAAA AATCAATATA AAAAGCCCTC	6720
GAAAGGGCTA ATAAATCTAT AAAATCAATA GGCGAGTAAC TAGCACAAGT GGACGTGCTT	6780

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TTTTATTGAC TATTACCACG ATACCACGCT TAATCTTAGG CTTGAACCTT CTTATCTGCA	6840
ATAGCGTCTG TCAAAGCTG AGAAAAGTTA AGCCCCATT CTCGTCCAA CTTATCTGCC	6900
CATTTGGTA TGGTCAAAGT CTTTTAATG GGTCCTGAC TTCCTAGGTA TTCTGATACA	6960
TCAACAGATA CCATAGAAAT AAAAGATTTA TCAAGGTCAT AGGTTGACAC GAAATCTCA	7020
TCATCTTAA AAGGATCATT ATCAATTAAA GACAAGCTAT TGATATCTGA TGGCTGAGGT	7080
AACTCTCCAT CACTCTCTAT CAAATCTGCA ACAGTTATCC CTAGCCACTC CGACCCCATA	7140
GCCAAAGCCT CAGAAATCCC CTCTCCTTGT GTAGCTGAGT ATTCAAAATC TGGGAAATGG	7200
ACAAAATAAG TCGCTTCTGT TCCGCTGTG TCGTCATAAT AAAATAAACG TGGATACGTA	7260
ACTAACATTT CACTACCTCC ATATCAAAAA GCAGGGACTG AATTTTACAA CCCAGCTTGC	7320
TTTCTTATCC CTCTTCAGT GTACTTATTC AGCTCACCAT GAAGGATTGT GATAGGTCTT	7380
TCCCCTTGCT TTTCCATTAA AATATGGGAG CCTTTACCGC CTCTAGTCTT TATCCAACCA	7440
TGGGCCGTA GGAGTTAAC CATCTCTTT TGTGTCTAG GCATAGCGCT TTTACCTCCT	7500
GACAACACCA TTATAACACG TGTTACACGT ATTGTAAAGG AGTGATACTT ATTATTCTAT	7560
TATACATAAA AGCCCTAGA TGTGGTTCTA AGGGAAACCA ATTTATTCTAT ACCTATTTTT	7620
CTAATGAGTA GTAAAAACTG CTTCTTATC GAGCAATTCA TCATCTGTAT AGTCAATTGT	7680
AAAAGTATCT CGATCTAAGA CAGATTGAGG CGGAGTTGAA TGAATCATAG GAACACTGCG	7740
TACTCTATAT TTTTATCTC CAATTTTAC AAACTGATAC TCTTCGAAAA TCAAATTCAA	7800
ACCACGTCAA CGTCGCCTTA CCGTACTCAA GTACAGCCTG CGGCTAGTTT CCTAGTTGC	7860
TCTTGATTT TCATTGAGTA TGATTAACTC TCAAGTCTTC GAAATCAGGA TTTCAACAG	7920
TTATTACAAG GAGGCGATTT ACTACTCAA AAACATCAAT TATTCTATTT TTCATATTTT	7980
TTCAACCCAT TATTAGAATG AACTCTTGG TAAGCAAAT CAAGTTAGA TTTAATGTTT	8040
TCGTACAAAT CTAAAATCTC TTTGGAGTA TCTTCCCGGA AGAAAAGTTT TCTTTCCCT	8100
GAAATAACTT GATCACTAAG AATCCAATGA CGAATTGTT TTGTAAAAAT CAAAATTCC	8160
TGACTTGGTA GTTCCATCAT TTCCATTGCT TATCACCTCT CTTTCATTA TAGTCATAC	8220
AATGACATTC AGCAATATTA TTTCTCAAGT CAGCACTTCC ACTTCTTTAG GCTCAACTAT	8280
CCTATTTGA GCTTTAAGGA AAATCAAATC TCTCATGCTG ATACCTCTCC TCATTAATT	8340
AAATAGTAAA AAAGATTCTA TCTCACTCCC TGATTATTAC AAAACCATTG AAATATCACA	8400
ACTAATAGGC TAGAATGGAC ATAGTAAGAT ATAGTAGATG AGTCATTCTA CTCAAATCCA	8460
CGTTAGAAAG GACTGCTATG CCAGACAATC TCGCCGTTCG CATGCGCCn GG	8512

(2) INFORMATION FOR SEQ ID NO: 130:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 2869 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 130:

CTCGTTCAA GGTTGAGTCT CTTGCAAATC TTGTTCGCGT TCTTCCTTTT GCCAAGGCAT	60
CTCTCCCATG GTGGTGccA GCCATTGTTG GAATCTTGCT CTCATTGGTT CTACCAAACA	120
AGCAAGAAAG CGATGTTTTT GAAATGGAAT AATCACTTAA ATCACTTTG TAGCCAAGTC	180
TACAGGAGTG ATTktCTTTT TTTATCCGAT GATAAATGTG TTATAATAGG TAGCGAAAGA	240
GGTGAAGAAA TGAATCAAAC AGTAGAATAT ATCAAAGAAC TGACACCCAT TGCGtCGCCA	300
ACAGGCTTTA CTCGTGAGAT TGCGGACTAT TTAGTCAAGA CTCTAGAAGG TTTTGGTTAC	360
CAGCCGGTTC GCACATCAA GGGCGGTGTC AATGTAAC TAATAAGGTCA AAATGATGAG	420
CAACATCGCT ATGTGACTGC CCATGTAGAT ACGCTTGGTG CTATTGTCCG TGCTGTCAAA	480
CCAGACGGCC GTCTCAAAAT GGACCGTATC GGTGGCTTTC CTTGGAACAT GATTGAAGGA	540
GAAAAGTGTAA CCATTCAATGT GGCTAGCACA GGTGAAAAAG TATCAGGAAC CATCCTCATC	600
CACCAAACTT CTTGCCATGT CTATAAGGAT GCAGGAACTG CAGAACGCAC GCAAGACAAT	660
ATGGAAGTGC GTTGGACGC CAAAGTAAC AGTGAAAAAG AAACCTGTC TCTTGGCATT	720
GAGGTCGGTG ATTTTATCAG TTTTGACCCA CGAACTGTCG TGACAGAGAC AGGTTTTATC	780
AAGTCTCGCC ATTTGGATGA CAAGGTCAGT GCGCGATTT TGCTCAATCT CCTTCGCATT	840
TATAAGGAAG AGAAGATTGA ATTGCCCGTA ACAACTCATT TTGCTTTTC AGTCTTGAA	900
GAAGTGGGAC ACGGTGAAA CTCTAACATT CCTGCTCAGG TAGTAGAATA TCTGGCTGTG	960
GATATGGGAG CCATGGGAGA TGACCAGCAA ACAGACGAAT ATACAGTGTG TATCTGTGTC	1020
AAGGATGCTT CTGGACCTTA TCACTATGAC TTCCGTCAAC ATTTGGTGGC TTTGGCGAAA	1080
GAGCAAGATA TTCCATTAA GCTGGATATC TATCCATTAT ATGGTTCGGA CGCTTCAGCG	1140
GCTATGTCTG CAGGGGCAGA AGTCAAACAC GCCCTTCTCG GTGCTGGTAT AGAGTCTAGC	1200
CATTCCCTATG AGCGTACCCA TATTGACTCG GTGATCGCAA CAGAACGAAT GGTCGATGCT	1260
TATCTTAAGA GCACGTTGGT GGACTAATAT GTGCCTTATT TGTCAGAGAA TTGACCTCAT	1320
CAAGAAGGAA GAAAATCCTT ACTTTGTCAA AGAGTTGGAA ACAGGCTATC TTGTGGTTGG	1380
AGACCACCAAG TATTTGAAG GCTATAGTCT CTTCTAGCC AAGGAGCATG TCAGCGAATT	1440

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GCACCATTG AAAAAGGAGA CAAGACTCCG TTTTCTAGAA GAAATGAGTT TAGTCCAAGA	1500
GGCAGTTGCC AAGGCCTTG CTGCTGAGAA AATGAATATC GAACTGCTAG GAAATGGCGA	1560
TGCTCATCTT CATTGGCATC TGTTCCACG ACGGACAGGT GATATGAATG GTCATGGTCT	1620
CAAGGGTCGT GGACCAGTCT GGTGGGTTCC CTTTGAAGAA ATGACAGCAG AAACCTGCCA	1680
AGCAAAACCG GATGAGATTA AAAGATTAGT CAAACGTTA TCGTCAGAAG TAGATAAACT	1740
ATTAGAAAATA AAGGAGTAGA AATGAAGAAA AGATACCTAG TCTTGACAGC TTTGCTAGCC	1800
TTGAGTCTAG CAGCTTGTTC ACAAGAAAAA ACAAAAAATG AAGATGGAGA AACTAAGACA	1860
GAACAGACAG CCAAAGCTGA TGGAACAGTC GGTAGTAAGT CTCAAGGAGC TGCCCAGAAG	1920
AAAGCAGAAG TGGTCAATAA AGGTGATTAC TACAGCATTC AAGGGAAATA CGATGAAATC	1980
ATCGTAGCCA ACAAACACTA TCCATTGTCT AAAGACTATA ATCCAGGGGA AAATCCAACA	2040
GCCAAGGCAG AGTTGGTCAA ACTCATAAA GCGATGCAAG AGGCAGGTTT CCCTATTAGT	2100
GATCATTACA GTGGTTTAG AAGTTATGAA ACTCAGACCA AGCTCTATCA AGATTATGTC	2160
AACCAAGATG GAAAGGCAGC AGCTGACCGT TACTCTGCC GTCCTGGCTA TAGCGAACAC	2220
CAGACAGGCT TGGCCTTTGA TGTGATTGGG ACTGATGGTG ATTTGGTGAC AGAAGAAAAA	2280
GCAGCCCCAAT GGCTCTGGGA TCATGCAGCT GATTATGGCT TTGTTGTCGG TTATCTCAAA	2340
GGCAAGGAAA AGGAAACAGG CTATATGGCT GAAGAATGGC ACCTGCGTTA TGTAGGAAAA	2400
GAAGCTAAAG AAATTGCTGC AAGTGGTCTC AGTTTGGAAAG AATACTATGG CTTTGAAAGC	2460
GGAGACTACG TCGATTAATA CTCTTCGAAA ATCTCTCAA ACCACGTCAG CGTCGCCCTTA	2520
CCTACTGACT GCGTCGGTTC TATTACAAC CTCAAAACAG TGTTTTGAGT cGATTCGTCA	2580
GTTTTATCTG CAACCTCAAA GCTGTACTTT GAGCAstGCG GCTAGCTTCC TAGTTTGCTC	2640
TTTGATTTTC ATTGAGTACA AAAAGTAAAC TTTTCTCTTG CAATTCCAGA TAAATAGTGT	2700
ATAATGGATG GGTATGTGAA AAACATACTT GTGGGAGGTA AAAATCTCTA ATTACCGCCA	2760
AAACCACAAA GGAGGATTTA AAAATGGCTA AAAAAGTCGA AAAACTTGTAA ATTGCAAA	2820
TCCCTGCTGG TAAAGCTACA CCAGCTCCAC CGGTTGGACC TGCTCTTGG	2869

(2) INFORMATION FOR SEQ ID NO: 131:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6186 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 131:

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CTGAATCCCT TATAGGAGTC CAGTAACCTT TTAGCCTCTA CTTTGCCCTC ATAGGCAGCT	60
TCAACATCAT TAAAAAAAAGA ArGCACTGAA GCAAGTTCTT CAGTGCTCCA CGACAAATCT	120
AGTGGGTAAC TATACTGTTT GTTCATTAAC TAATACCAGC TCTCATTCTT GCTTCTTTA	180
GTTCTTGCTT ACGATAACTA CGAGGGAGAA AAGCACGAAT CTCATCTTCA TTAAAACCGA	240
TTTGCATACG CTTGGCATCA ATAATAATTG GACGACGAA AAGACTAGGA TACTGCTCAA	300
TCAAATGAAG CAATTCCGAT ACCGAAATAC TCTCTACATC AATATTCAAT TTTTGAAAAAA	360
TTTTGAAACG AGTTGAAATG ATGTCATCAG TACCATTTC GGTCAAGGAA AGGATGTGTT	420
GCAATTCTTT TCTTGTAAA GGACTGGTCA TAATATTGTG TTCCACAAAG GGAACCTTATG	480
TTTTTCTAAC CAGGCCTTAG CCTTACGACA TGATGTACAG CTCGGTGATA GAAATAGTGT	540
AATCATGCTT TTCTCTTCTT ATCTATACTT TGCTACTTCT ATTATACAAA AAAATAAAGC	600
GCTTGACTAG GGATTTTAG AAAAAAAGCC TATTTTTCA AGAAAAATAG GCTTTTGCG	660
AACGATTGAC ACAATTGGAT TTGGTTAATT CACTCTTAAC GATGGTTTA AACGATATAT	720
ATTTTTATAT ATGTAAATTA AAAACATCTT TCCTTTCACT TCCTACGACT TTTCAGATAC	780
AGATAGCCAA AGAAGTTTC ATAGAGGGCA AAAAAGAGGA GGAAGGCATG AAGAAAGAAG	840
GTCTCTGGCA AAATCATAAT AACAGGATCC TTGGCTGGAT CAAAAAGCCA GGTATCATCT	900
CCCACAAAGA GAATTTGATC GAAAAGAGTA AAAAAGAGGT CAAAACCAAT CAAAACCTCC	960
CCAAGTCCAA TCATCACAGG TAAGACTACT AGAGCCAGGA GACTTTTCG ATAAAGAGAC	1020
AAAAAGTCCT TTTTCACAAT CCTATTGACA AAGACATAGA AACTTGGCAG TGTCACTAGA	1080
GCTACTAGCT GAACCAAATG AAAGAGATTG TTGACCACTG CGAAATGGTG CAGACCAGCT	1140
GCTGACGAAC GAAAATCAGG CATCTGTAAG ACCTGACTAA AAGGATTGGT CAGATAATTG	1200
ATCAAGATAT GAAAATTGTA TTGAATGGTT TCTGGTTTTA GATAGACTCG ATTGTTAAG	1260
TTTAGCCACT GAATCTCCAT AGGATAGAAA ATCCAAGCCA GATAAATGGT CAGAAGGATG	1320
GAGAGGGAGA GGAGAAAGAG CATAGAGCCC CAAAAGATCA ATTTAGTTT CATCAAAATC	1380
CCACTCCGCA AGGCTAGAAA CCACATGTGT CGGTGCGATT GGCAGGCCAG CTACTTCTTC	1440
TGCCTTAGTA AAACCTGTGTC TCACCAAGAG CGTTGGAATG CCATTGTCAA TCCCAGCCCG	1500
AATATCAGTC AAATAATTGT CCCCAACCAT GATTAACCTCT TCACGTTCCA AACCTAAGTG	1560
CTCAACCGCC TTGTCCATAA TGATGGCATT TGTTTTCCG ATATAAACCG GCTTCACTCG	1620
TGTCGCTACT TCAAGCAGCG TAATCAGTGA GCCAGCACCT GGCAAAAGAC CGCGTTCCGT	1680
CGGGATGTTG AGGTCAAGGAT TGGTTCCGAT AAAATGGCA CCCTTTGAA TAGCAAGAGT	1740

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TGCTGTGGCA	AATTTTCAT	AGTCGACTTG	CCAATCCAGA	CCAACTACCA	CGTAGGCAGG	1800
TTTTTCCCTTG	TCTTCCACAT	AACCAGCCGC	CTTGATGGCT	TCCTTGAGTC	CTGCTTCTCC	1860
GACGACATAG	ACGGTCTTT	CAAGCCCCAA	ATCATTCTATA	TAGTCGATGG	TTGCCAAAGT	1920
CGCTGTGTAG	ACAGTCGATA	GGGGCGTATC	GATATTAAAA	TTCTGAGGCCA	ACATCTCCTT	1980
AACACTCTCT	GGAGTGCAGG	TTGTATTGTT	GGTTACAAAG	AGATAGGGAA	TGTCCCGCTT	2040
TTGCAATTCA	TGAACAAAAG	TCTCTCCAGC	AGGGATTCCGG	TCTTCCCCT	TATAAATGGT	2100
TCCGTCTAAA	TCAATTAAAT	AGCCTTTATA	TTTCATCTAT	TTCTCCCTAA	GCCTTTTTTA	2160
TTTCTTGCCA	AGTAATGATT	GCTTGGCAT	TGATAACCCC	ATCACTTGTA	ATTCATGCT	2220
TGCTTTCCAG	TCCAGTCCGT	TCAACAGCCG	ATGTAATCAC	CCCACCTGGT	CGAACCTCCT	2280
TGACATACTT	GAGGTTGATT	TTCTTGGAA	TATAGTGGGT	CAAAAAATCC	GCTCCCATGA	2340
CCTCAAAAT	CCAGTCCAAG	TATTTACTGT	TATTGACATG	ACCATTCTATA	TCCAAGTCGT	2400
AAAAACGAAC	ATGGTAATCC	TTGCTGATCG	GTTCTTCAA	GGACTCATAC	TTCGGTCCAC	2460
GGATAAGTTT	TTTATCAAAA	TCAGACTGGT	AAGGAGCCAC	AATCTCAGGT	TCAACAACAT	2520
GGACTTTTCG	ACTGTGGCGG	TCCATGAGAA	CAAAGGTCGC	CATCATGTGG	ATGAGCTCCT	2580
GCTCCGCTTC	ATTATAAAATA	GTAAAGCGAC	GGTAGCAAA	AAGTCGATTG	TAGCTCAAGG	2640
CTTCCGTTTC	GATGGTAATT	TCTTCCGCAA	AACGAGGCAA	ACGAACCACC	TCAATATCAT	2700
ATTCTACGAT	AATCCAGACC	AGATTATATT	CTTCCAAAAT	GGCCTTATCA	CTAACTCCCA	2760
GTTCAATCGA	CTGCATCCCT	GAAACTTGCA	GTGACAGCAA	AATCACATCT	GGAAGTTTGA	2820
TATGACCGTT	CATATCAGCC	ATATCAAAG	GAATTTCAT	TTTCATTGA	TAAGTTAACG	2880
CCATGATCCT	ACTCCAAAAT	AAATCGTTCT	GCTACAGTAT	CTCCCAAAAA	GAGACCTCTC	2940
TTTGTATGC	GAACGTGGTC	ACCCTCAATC	TGCATGAGGC	CTTGTGAAAC	CAAATCTCTG	3000
ACAATTCTC	CATAAAGTCC	AGCAAAAGAC	TGTCAAATT	TTTCCTCAA	TCGCGCCATG	3060
GAAACCCCGG	ATTCTTGCG	GAGTCCCAAG	AACATTCTT	CTTCCATTG	CTCCTTTGA	3120
CTCAGGTGAT	CTTCTGTAAT	ACAAGCATTG	CCTTCCCTCAA	CCGCACTGAG	ATAATGACGA	3180
ATGGGACCAT	GATTTTATA	GCGTACTCCA	TTGACATAAC	CAGATGCC	TGCACCAATA	3240
CCATAGTATT	CAGCATTGTC	CCAGTACATG	AGATTATGAC	GACTTTCAA	ACCGGGTTTG	3300
GAGAAATTAG	AAATCTCATA	ATGCTAAAA	CCCGCTCGCT	CCAGCTCTGC	AATGATGTAC	3360
TCAAACATCT	CCGCTTCTAG	TTCCCTCTTA	GGCAGAGGCA	ATTTCCCACG	TCGCATCCGG	3420
TTCATAAAAGA	CCGTATGGTT	TTCTAAATC	AAACTATACA	AACTCATGTG	GGGAATATCC	3480
AATCCAATGG	CTTTAGGCCAC	ATTTCTTTT	ACTTGCTCCA	TGGTCTGACC	AGGCAGAGCA	3540

899

TAAATCAAAT	CAATGGAGAT	ATTGTCAAAA	CCAGGCCAGTT	TCAGGCGATC	GATATTTCA	3600
TAAATATCCT	TCTCCAAATG	ACTGCGCCA	ATCTTTTCA	ACATCTTATC	ATCAAAGGTC	3660
TGGACACCTA	GCGAAACACG	ATTGACAGCC	GAATTTTCA	AAACAGCTAT	CTTATCCGCA	3720
TCCAATCGC	CTGGATTGGC	TTCAATGGTC	AACTCTTCCA	AGACAGACAA	ATCCAAGTTT	3780
TTAGTCAAGC	CATTCAGTAA	CACCTCCAGT	TGCGGAGCCG	ACAGGGCTGT	CGgTGTTCCA	3840
CCACCGATAT	AAAGGGTTGA	CAACTTTCA	ATATCATAAG	AACGAAACTC	TTCCAGCAGA	3900
TGCTCTAAAT	AGCTGTCGAC	TGGCTGATTT	TTGATGAAGA	CCTTGAAAA	ATCACAATAA	3960
TAACAAATCT	GGGTACAAAAA	TGGGATGTGC	ACATAGGCTG	ACGTTGGTTT	TTTCTGCATA	4020
GTAATTATTA	TACCACAAAG	ACTAGATTCC	AGATAAAAAT	CACCATCCCC	AGATACATAG	4080
TCCGTCCGGA	GATGGTGATG	GTtttATTCTT	CTGTTATATC	AATCACAATC	TCTTCTGAGT	4140
CATCAAGAGC	TTCGGCTTT	TCTTGCCATT	GCTCCTTGAG	ATTATTTAAT	TGATTTTTG	4200
ATGCTTCTGT	CGCTTGAAAA	GCATAGGATT	TAGTTGAGC	AACTATACTG	TCCACAGTGA	4260
TTTCACCTGA	CTCAACCTGT	TCTTTGTTT	TCAGAACAAA	ATCTGTAGCC	TGCTCCTTAA	4320
CTTCTGTCAG	TTTTTACAG	ACTTGCTCCT	TGGCATACTC	CGGATCTTCT	CTCAAATCAT	4380
CTAGAAAATC	TTGAGCCTGA	CTGCAAACCT	GTTGCCCTT	ATCACTTGT	AAAAACAAGG	4440
CAAGAGCTGC	ACCTGAAACG	GTTCTAAAAA	GGATTGAGGA	TAATTTACCC	ATAAGGATTG	4500
TCCTTTTTA	TTTTTGAAA	AATTACTTG	CAAGACGAAG	AGCTGACAGA	CTTGCACCAG	4560
TCTTGAGTGT	TTTGAAACCA	GCTGATGAAG	CTTTCTTGCT	CAAGACACGC	GCATGGTCAT	4620
TGAGGTCTGA	AACAGATAGA	GATAATCTG	CAACAGCACT	GAAGAGTGG	TCAATCGTAG	4680
CCACCTTGAC	ATTGATATCA	TCTGCCAAGA	CATTGACCTT	AGCCAACAAAC	TCATTGGTGT	4740
GATGCAAGGT	CACATCCACA	TCTGAAGTCA	AGGTTTAAT	CGTCTTTCT	GTTTCATCGA	4800
TGACACGACC	AAGCTTTGT	ACAGATAATGA	TCAGATAGAC	CAAAAGACA	ATCAAAGCTA	4860
GGGCAACAAG	AATATATGCA	ACTTCTAACCA	TTTAGTTTTC	CTCCTCTGTA	ATATAGTAAG	4920
GGGCCTTCTT	TCGATTTGAA	AAAATAACGA	TCATTATACC	GAGACCGATA	AGGACAAC	4980
ACAGCCATTG	GGACACTCGA	AAGCCGAAGA	ACATGAGACT	ATCTGTTCGC	ATACCTTCGA	5040
TAACCATAACG	ACCGAAACCA	TACCAAATCA	AGTAAAAGGC	CGTGATATGA	CCTCGTCTGA	5100
GAATCTTCCA	TTTCCGTCTA	AAAATCAGAA	TCAAGGAAA	GCCAAGCAGA	TTCCATAGAG	5160
ACTCATAAAG	GAAAGTCGGT	TGACGGTAGC	TCCCCTCAAT	ATACATCTGG	TCACGGATAA	5220
AGCCAGGTAG	ATAATCCAGA	TTATCCACTG	TTGCACCATA	AGCTTCTGG	TTAAAGAAAT	5280

900	
TACCCCAACG CCCCAAAC TT TGAGCAATCA TAACGCTAGG CGCCGCAATA TCTAGAAAAT	5340
CCCAAGTATT GATGAGTTA CGGTCAAGCAA AGATATAGAG CACAAGAGCC CCAGTTATCA	5400
AACCACCGTA AATGGCCAAA CCACCATTCC AAATGGCAAA AATCTCTCCT AAATTCTGAC	5460
TATAGTAATC AAATCGGAAA ATAACATAGT AGAGACGAGC TCCTAAAATA GCCAAGGGAA	5520
AGGCTACTAA GATAAAATCT AAAATATCGT CTGGTATGAT CTTCTTCCTA GGTGCTTCCT	5580
TCATGGTCAA ATAAACCGCA AGAACATCAAGC CTGTCACAAT ACATAAGGCA TACCAACGAA	5640
TGGCTAGGGG TCCTAGTTGA ATAGCAATTG GATCAAGCAT TTTGCACCTC ATTCGAGCG	5700
ATTAGACTTG TCAGTCGTTG GTCGAACAAA CGGGTCGCAT CAAAGCCCCT TTCCTGGCA	5760
CGATAATTCA TGGCAGCTGC CTCAATCACA ACAGAGATAT TACGACCTGT TTTAACTGGA	5820
ATACGAATAC GAGGAATGtA CGCCAGAAAC TTCAAGTTCC TCTGCATTAT TTCCAAGACG	5880
ATCAAAGGTC TTATGCGTAT CGTAATTTC CAAATAGACA GCAAGCTGAA CCTGTGAAGA	5940
ATCCTTGACA GCACTCGCAC CGTAGAGACT CATAACATCG ATAATACCAA CCCCACGAAT	6000
TTCAATCAAG TGTTCAAAAA TTTCAGCTGG TTCACCCCCAG AGAGTAATCT CATCCTGGC	6060
AAAGATATCG ACACGGTCAT CGGCTACCAA ACGGTGACCA CGTTTGACAA GCTCAAGACC	6120
TGTCTCGCTC TTACCAATTC CACTATCTCC CTGAATCAAG ACGCCCATCC CATAAATATC	6180
CATCAA	6186

(2) INFORMATION FOR SEQ ID NO: 132:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9541 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 132:

GAAAATCACA ACCCTTTTG CAAAATTTT GAGATTATTT TCACAAAC TT GATTTTCAA	60
AGTATACTCA ATAAAAATTA AAAAATCCA CTACGTCAAG GCGAGGCTAA TGTGGTTTGA	120
AGAAATTTTC GAAGAGCGTG AATGAGTATC ATCTATAGTA AAATAAAAAAA ACTGAACAAT	180
TTGGTTGGGG ACAGCCAAAC CAATTCTCA CAATGTTCA GAAACAAGGG TGTGCTATT	240
CAATTTCAGC CTACTATAAC TGTCTAGAT TGCTGAAACA AAGTCTAGGT AAAAGTCTTC	300
ATAATAAAAA GACCTCCTAT CAAGTGTCA AAAACTTTGA TAGGAGGTCT TGTGCTATT	360
AAATATTTAT CAAATTTCT ATACAAGTGA GCTGTTAGCC AGGTTCTTC TATTCTTC	420
ATTTCAATGA ATGGATTTTT TACTAATCT CATAACTGGG AATTTGTCTG TGTAAAAATA	480

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GCGAGATAGA TGGTATTAT AAAACACTCA AGACAGCTAG ACTAATATCA TTTAAAACAT	540
TATCTTCTTT TGAGCGACTG TTGGTTACCA ACATAGCTAA ATTTCCTGCA TTTTCAAATT	600
GATAGGGTTC TGATTTAGCA TTCACAACCA CCAAGAGGTG TTCTTGCCG TGAACCTTCAT	660
AGATAAGGTA GCGCTATGT TCAATCGCAG AATGCACAAA GACATGATGG TAAATTCAT	720
CATAGCTAGA GTAAGAAAAG GCACCAGTTT TTGTCTTCAA TCGGATGACT TGACGGATAA	780
ACTCAAACTACT GTCTTGACGC TCATTAATCA AGTTCAGTT CACTTGGTTC ACACTGTCAG	840
GAGCATTATA GCTATTTCATC GCACGCTCTC TATCATCATG GGTCAACTCA CCATTTTCAC	900
CAGTCGCAAC CAGTTGGTA CGACCAAATT CTTGACCGAT TTCCATAAAG GCCATCCCC	960
GCATGAGCAG ATTCAATGGCT GTGGCAGTTT CGACCTTGCG CATGATTGTC TCTGAACCTT	1020
GGTCTGGATG AAGGGTTGCC AATAAAATCGT GAAGATTGTA ATTGTCAATGG GCTTCTACAT	1080
AGTTAACGAC CTGATTGGA TGTGTATAGC TTCCATAATTC ACGACTTCCT AGGATTGCTT	1140
TAGCTAGAAT TGGCTCTGTC GCAGCACCAC TGACAAAACC TGACTTGATA GCACCATAAA	1200
CTTCTCCCCC TTTGACAGCA TCGCGCTGAT TGTCAATAAA GAAACCAATA TTTGGCATCT	1260
GGTAGGCATT GTCCTTCTTG GCCTTATCAT AAGGGCAAG ACCTGTTCCC ATATCCATC	1320
CTTCTCCATA GAGGATAATG TTGGAGTCGA TTTCATCCAA GCTTTGACGA ATCATCTGCA	1380
TGGTCTTGAC ATCATGAATC CCCATCAAGT CAAACCGAA GCCGTCAATA TTATATTCC	1440
GCACCCAGTA TAGAAGAGAA TCAATCATAT ACTTGCAGAA CATTCTGTGT TCACTGGCTG	1500
TTTCATTTCC AACACCCGTT CCATTCTGGA AGGTACCATC TGGATTCTATA CGATAATAGT	1560
AATCAGGGAC TGTTGTTGG AATGGTGCAT CAACAACTGA GAAGGTATGG TTATAGACTA	1620
CATCCATAAT GACTCCAATA CCCGCATCGT GATAAGCTTG AACCATCACC TTCAAATCAC	1680
GAATGACCTG AGCTGGATCA TCTGGATTAG TTGAAAAACT AGTTTCTGGC GCGTTATAGT	1740
TTTGTGGATC ATAACCCAG TTGTAGGTTA CATTCCATC CTCATCGTAT TCTTTATGAC	1800
GGTCTGCAAT TGGTTGCAAT TGAACATAAT TGTAGCCAG CTTCTTGATG TAATCAAAG	1860
CAGTTGACTG GCCGTATTGG TTAACTGTTG CAGCCTGAGC AGCACCCAAG AAAGTTCC	1920
GAAGATGTTG ATCTACACCC GATGTAGGTG ATTTAGTCAA ATCACGAATG TGCATTTCAC	1980
AGATAACTGC CTTACATGGA TTTTCCAAGC GCCAAGTAGC CTCCGAACCG TGCTTAACCT	2040
CGAACGTTTC AACTGCTTT TCTACATGGC TCAGAATAGC TGAACGTTTG CCATCAGGGC	2100
TGGTCGCGAT TGTATAAGGA TCACGTGTCA GTGTTGGTG ATGAGGGAAT TGGACTTGAT	2160
ACTGATAAGT CTTACCTACC AAATCTTCTT CAACATCCAA ACTCCAGACCA CCGATTGTAT	2220

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TGTCCTTATG ATTATAAGAG TAGCTATTGC CTCTTTCAT CTCAAAAGTC TTCCAAACGG	2280
GTGCATCATT AGCAGCTGAT TCATAAACGA CAACTTGCAC TTCTGTCGCT GTAGGTGACC	2340
AGAGAGAAAA ATGAGCCTGA TTGTCCTCTA CACGGCAACC CAATTCTCCT TGGTAACCCC	2400
AATGATGATC AAAACTAGCA CTGTTAATGG CCTTATCAA GGCAAAAGGA TTTTGATTTT	2460
TATAGAAAGG ACTGGCAATA GCAGGGATTT CAGAGTAATA AATCCTATCA TCGCCTCCA	2520
AAATCCAGAC CTCTGTTAAT AGGGGATAGT GATTAAAACG GATAGAATAT TCTTTACTAG	2580
TTTGACCTGT ATGAACCACA AAATTCAAGC TTTCTATAAC ATGTGAACCTT GGGTGTCAA	2640
AGCTAAATAA AGCTCCAAAA TAATCTTCTT TGTAGGTTAG CAAATCAATT CGTTGATCCT	2700
GACTTTTAC AAAGGAGCAA GTGTCATATT CTCCATTCTT ACGATGGTAA TGAATGCGCA	2760
TAGGGTAGTT ATACATTTT TATTTTTCCT TTTTACTTTG TTTCTATTTC ACTAATAAAT	2820
TTTTGTCAAT CTCGCTCAA TTAACAGACA TAGTCATATT CTCTAAACTC TGTTTTAAA	2880
CGATCCATTA CAAACTTTCT AGCCATGCCT CATCTCTGAC CTGGATACCA AGTTCTTGTG	2940
CTTTTGCAG TTTACTTCCA GCGTCTGCAC CTACCACGAC GAGGTCGGTC TTTTAGAAA	3000
TACTACCTGT CACTTGGCA CCCAGACTTT CGAGTTTACT TTTAGCTCT GAGCGCTTGA	3060
GTCGTTCAA TTTTCTGTC AATACCACGG TCAAACCTGA CAAGGCCGCA TCCGCTACTA	3120
CCGTCTGTCC TTTATAGTCC AGATTGACCC CAGTTCTTT CAATTCTCTG AGCAGAAATT	3180
CAGAGCCTTC TGTCGCAAAA TAAGTCTGAA GACTTTGGC AATCACGCCA CCTAGACTTT	3240
CAATACTAGC CACTTCCCT GAATCTGCCT GAGACAGATT TTCAATTGAA TGGAAATATT	3300
GAAGTAAAAG CTGACTAACCT TTGCTTCCGA CATGACGAAT TCCCAAACCA AATAAGAGCT	3360
TCTCGGCAGA ATTTCCCTT GATGCTTGGA TAGCCTGATA CAGTTTAGCA GCGGACTTT	3420
CCTTAACTCC CTCTAAAGG AGGAAATCCT CTTCTTGCAA ACGATAAATA TCCGCCACAT	3480
CCTTGACTAA ATTAGCAGCA AAAAGCTTCT CAACAATAGA TGGACCAAGG CCTGTAATAT	3540
TCATAGCATC ACGAGAAGCA AAGTGAATCA AGCCTTCCAT GATTGAGCA GGGCAACGCG	3600
GATTGATACA ACGTAGGGCC ACTTCATCTT CAAAGTGCAGA CAAGTCAGAG TTACAACCTG	3660
GACAGTTTGT AGGGATATCT AGTTTTCTT CAGAAACCCG TTTGGACTCT ACCACACGTA	3720
AAACGGCAGG GATGATGTCA CCAGCCTTAT ATACAATGAC CGTATCGTCT TTTCGGATAT	3780
CTTTTCAGC AATATAATCT ACATTGTGCA GGGTCGCACG GCTAACAGTC GTACCGCAA	3840
GTTGTACTGG TGTTAGATTA GCAGTTGGAG TTACAACACC GGTACGGCCA ACTGTCCAGT	3900
CAACTGATAA GAGTTGAGCT TCTTTTCTT CGGCAGGGAA CTTGTAGGCT ACTGCCACT	3960
TTGGAGCCTT AACTGTAAAA CCAAGTTCTT CTTGACTTGC TAGGTCGTTG ACCTTGATTA	4020

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CCACTCCATC AATATCGTAA	'GGCAGATTTT CCCGTTCTG TCCTACTTCT	TGGATAAAAT	4080			
TCCAGATTTC ATCTATGTTT	TCAGCCAAGA TTCGCTTAGG ATTGACCACA	AAACCTAGTT	4140			
GTTCTAGGTA CTTCAAACCC	TTTCTTGGC TATCACGAGT TGAAGGGCTG	GCTTCTTGAT	4200			
AGAGAAACGT TGCAAGATTA	CGCTTGGCAA CTACTGCTGT ATCCAATG	CGCAGAGTTC	4260			
CTGCTGCCGC ATTACGAGGA	TTAGCAAATT CAGGCTCTCC	ATTTCTTGG CGCGCTTGGT	4320			
TAACTTGGTC AAAGGAAGCG	CGTGGCATGT AACATCCCC	ACGAACATGTG ATATCTAGTT	4380			
CTTCTGGCAA AGTCAAAGGG	ATGTCCTTAA CACGCTTGAG	GTTTCTGTG ATATTTCAC	4440			
CAATTGAACC ATCTCACGT	GTTACCCCAG CAACCAAAT	CCCCTTTCA TAAGTCAGCG	4500			
AGATAGATAA GCCATCGATT	TTCAGCTCAC AAATATAGGT	CGGATGAGCC ACTTCCTTAC	4560			
GAACACGCCG ATCAAAGCA	TCTAGCTCCT	CACATGAAAA AGCATCCTGC	4620			
GAGGATACTG ATGACTGTAT	TTTCAAAAC CATCTAAAAC	CTTGCCACCA ACACGATGAG	4680			
TCGGACTGTC TGCTAGCACT	TGCTCTGGAT AAGCAGTTTC	TAACTCGACC AACTCACGGT	4740			
AAAGGCGGTC ATACTCACTG	TCTGAAACCG AGGGATTATC	GCTGGTATAG TACTCAGTCG	4800			
CATAGCGATT GAGCAAAGCG	ACTAACTCAT TCATTCTTT	ATTCTATAAGA CCATTTTACC	4860			
ATAAAACAAG CCCTCCTCAC	AAACGAGAAG GGCGAAAAAA	ACACTTAGTT TGAAATTATT	4920			
TTTGAAACTC AAGCAACCTT	ATATCAATTT	TTCAAAATGA GTTCGAACAT ATCCGAGAGC	4980			
TAAGAAATAT AAGGCTACAA	CTCCAAGTCC AATAATCAAG	AAAGAATAAA GATGGACACT	5040			
TGGCAAGACT GTCATAAAC	CTTTGCAAT AGGCATAAAAT	AGAATAGCTA AGGTAAAAAT	5100			
TGTACTCAGT ACTCTTCCAA	GAAATTGCT CTCAACCTTG	GTTTGTACTT GAGTAAAAAA	5160			
GTGAATATTA AAAATCGTCA	AAACAAATTC	ACAAACTAAA TTTCCAGAAA AGGAAAGAAA	5220			
AGTTGGAAGT GGTAACTCCA	TCATAAAAAC	TCCGACACCT GTCAAAGCCA	5280			
AAGATTATAA ATATTAGCTT	TAATTTACT	AGCTAGAAGA GCCCCAATGA	TGGAACCAAT	5340		
AGCCCCCATA GTTAAATAC	TTGCATAGGC	TCCTCTGAC CCGTAAAGCT	GATTGAAAAA	5400		
GGGAAGTAGA AATTCAAAG	CTGCAAAAAAA	GAAATTAAACG CTGGAAGCTA	CCAGCAAAAG	5460		
GAAGAAAATT TCTTGCTGAT	GCCAGATATA	GTGTAACCCA	TCCTTGATAT	CTACAAAAAT	5520	
ATCTCTCCA GTAAAAGCCT	TTTCTCTTG	AACTTTTGCT	TCCTCTTTG	GAAGGAAAGC	5580	
CACTAGAACAA	AAAGCAATGA	AAAAAGTCAG	CGAGTCTAGC	AGTAGCGTCA	TATGGAGACT	5640
TGCAAACGT AAAACAAGGA	AGGAAAGAAC	AGGAGAGCTA	ACACCTACAA	CCTGCAAAAC	5700	
CAGCTCTAAG CGAGAATTAT	AGATCACAAT	CTCATCTTC	TCCACCAC	TAGTTATGAT	5760	

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AGCTTTATTG GCTGTCCGAG AAAAGGCAAA AGCAATAGCC TGCACAATGT TAGCAACAAT	5820
CAAAGCGCCA ATCATCCAGC TATCATTCCCT TATGAAAGAA ATAGCCAGAC AAAGAATCCC	5880
ACAAAACAAGA TCTGCCGTCA TTAAAATCTT ACGACGAGAA AAACGGTCTG AAATAACTCC	5940
GCCAAAGGGA TTGACGAGAA TAGATGTGAC GAGCTCAGAA ATCTGATACA TTCCTAAAAC	6000
TGTCTGTCCT ATAGTCCCCA TAGAAGCCAA CCAGACACTA TTTCCATAAT CATAGAGCAT	6060
ATTTCCCATT TTATTGATAG CCCCACGGCT AATCAACTGC ACTGCATAGC GATTCAATTATT	6120
AAAGCTCCTC TCAAATTGG AAACATTGT ATCAAAACCG AAAGGAGCTT TTTATTTTTT	6180
CCCTTATTGG GGAAAATTAA CTTTGACAA ATTTTCGTA GTGTTCCCTGA TAATAGGCTA	6240
CTTGCTCTGG AAGACCTAAC ACATCAAAAA TATGCATGGC CTCTGCATC TGCTTACAGC	6300
CTTCTTTACA CTGTCCTTT TGATATAAGG CAAAACCTTT TAAATAATGG AAAACATTAC	6360
GCTCATAAAG CTTAACACCT TTGTCATAAA TCTTCTCTGT ATAAGCCTCA AAATAGTTGG	6420
CATTATAAAA AGAAGAATGC TCTAAACAAT GCTGGTAACA ATTGAGGGCC AAAATCAACA	6480
CTAATCTCTT ATGGCGACTA ATCTCTTGGT AAAATCCCTC CCTCTCCATA ACTTCTCTAC	6540
CAATCCGAGT GACATAGTCT ACATCGTAGA AACTATAGAG GTTACCGAAA AGAATCAACT	6600
CATACATGGT CCATTCTCT TGTTTGAAGA GATAATCTGC TACCTTACCC AAATCATCCT	6660
GCTTCATATC ATAACTCGCA TCTCTTGAC AAATCAGACC TTGTAGCAAATCCAGTTCA	6720
GCTCAAATA AAGGGGAGTC GTCGAACCT TAGACTTTTC AAGTTGTTCT CTTTGAAGCT	6780
TTTGAAAACC TGCAATATCG TTTGAATAGT AAAGTGGGAT AATCTGTGCC ATCATAGACA	6840
CATGTTCATG ATTATGAAAA TTCCCTGCCT TATCCATGAA ATTTTCGATT GTTACATGAA	6900
TGTTATCCAA AATCTCAAAG AAACGGGAGA CTGCCAGGTC AGACTCCCCA AGCTCAAAGC	6960
GAGATAACTG AGAGGTAGAG CAGGATTCGC CTGCTGCTTC CTTTAAAGAA TAATTCCAC	7020
TTGTTCGAAA TTCACGAAAT ACTTTCCAA GATGTTCCAT CTTTACACCT GCTCTGATAA	7080
TTCTTCCCAC TCAAGCATAG CTTCTCCTG ACGATGGCTG ATTTGTCCA GCTCAGCCTG	7140
TAATTCCATG AGTTTGTGG CATCGTTGT TTCCAACATT TGTCAGAAA TGGCTTGGCT	7200
TTGACTTTCT AGCTCTCAA TTTCACTTC TAGACTTTCG ATTTGTGCA TGAGTTGCG	7260
AACTTCTTT TGACTTTCTT TCTGGGCCTG ATAGTCATTG ACTGGACTTG CTTCTTTGC	7320
TTGATTGCTA GTTGAAGCTT CCTCAGTCTG ACTCATTCT GCTGTTGCTT TCTTCTCAAC	7380
ATAGTAGTCG TAATCTCAA GGTAGAGAGT TGAACCATT TCAGACAATT CCAAAACATG	7440
AGTTGCCACA CGATTGATAA AGTAACGATC ATGACTGACA AACAGCAAGG TTCCATCAA	7500
GTCAATCAAG GCATTTCTA GCACTTCCTT ACTATCAATA TCCAAGTGGT TGGTCGGCTC	7560

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ATCCAGAAC	TC AAAAGTTAT	TGTTTCCAT	AGACAATT	A GCTAAAAGCA	AACGAGCTTT	7620	
TTCGCCACCA	GATAGCATGC	CGACTGATTT	TTAACATCA	TCTCCTGAGA	AAAGGAAGGC	7680	
TCCAAGACGG	TTGCGGATTT	CAACTTCTGG	TGTCA GTTTG	AAATCATTCC	AGAGTCATC	7740	
CAGCACCGTA	TTACTTGGTG	TCAGCTTGCT	TTGGGTTGG	TCATAGTAAC	CAACCTCAAC	7800	
ATTAGCGCCA	AAGCGCTTT	CTCCCTTGAT	AAAAGGAATC	TGGTCCACAA	TAGACTTGAT	7860	
AAAGGTTGAC	TTGCCGATAC	CATTGGACC	AACGATAGCG	ACAGCATTCA	TCTTACGAAG	7920	
ATCTAGGTTA	ATCGGTTGTG	ACAAGACTTC	CCC GTCATAG	CCAACAGCTG	CATTTCAAC	7980	
AGTCAAAACA	ACATTGCCCG	ACGTTTTTC	AGACTGGAAG	GTCATGTTGG	CTGATTCTT	8040	
GCCAGCTTCA	GGCTTGTCCA	AACGTTCCAT	TTTTTCCAGT	TGTTTACGGC	GAGATTGAGC	8100	
ACGTTTAGTC	GTTGAAGCAC	GAACTAGATT	GCGATTGACA	AAGTCTTCCA	GAGCAGCGAT	8160	
TTCCTTCTGT	TGCTTTTCAT	AGTTTTTGC	CTCAGTA	ACT AGCTTTGCT	CCTTCAATT	8220	
GACAAAACGA	GAGTAATTCC	CCACATAGCG	ATCCAAGGAA	TGCTTGGTCA	AATCTAGCGT	8280	
AATTGTCGCA	ACCTTGTCCA	AGAAATAACG	GTCGTGGCTG	ACGATAATGA	GGGCACCGCT	8340	
ATAGTTTACC	AAGTAATTCT	CTAGCCAGGC	GATGGTTCA	ATATCCAAGT	GGTTAGTTGG	8400	
CTCGTCCAAG	ACCAAGAGAT	TGGGTTTTTC	AAGGAGCATT	TTGGCAAGTG	CCAAACGAGT	8460	
ATTTTGACCA	CCAGAAAGCT	CAGCAATT	CATCTGCCAC	ATAGACTCGT	CAAAC TTGAA	8520	
TCCATTCAA	ATCGCTCGAA	TATCAGCTTC	ATAGGTAAAG	CCACCTGCTT	GGC GAAAATT	8580	
CTCAGATAAG	CGGT CATAAT	CTGACATCAG	TTTATCCAAA	TCCTCACCA	AGCTTTCA	8640	
CATCTCCAGC	TCCATCTGAC	GCAGTTGTCT	CTCCGTCCGA	CGCAAATCAT	TAAAGACATG	8700	
AAGCATTCA	TCGTAGATGG	TATTTTCAGA	CTCAAAACGG	CTATCTGGG	CTAGGTAAGA	8760	
CAGAGAAATA	TCTTTTTCT	TATTGATTTC	TCCGCTAGTT	GGCTCCTT	CTCCA ACTAA	8820	
AATCTTCAA	AGAGTAGACT	TACCTGCACC	ATTTTCCC	ACAAGAGCAA	TCCGATCTCG	8880	
TTCATCAACC	TGCAGGGTGA	TATTATCGAA	AAGAACCTCT	CCTGCAAAG	AACGTTCAAT	8940	
TTTATTAGCT	TGTAAAATAA	TCATACAA	AGTATAGCAT	GTTTCCCTAA	GGCATTCAAG	9000	
ATAATCGTAA	GTCTTTAGT	ACAAC	TTTA	TAACATAAAA	TAAACTAAAT	TATGTATATT	9060
TTATATTAGA	TTACTTCACT	ATCTGTTGG	ATTTCTAAC	CAGCTAATCT	TGTTTCAAAT	9120	
AGTTATCGCA	CAAGTCTATT	ATTTAATTCT	TTTCATCATT	TACGTACGTA	TAGCAGATTG	9180	
AAATAAGATG	AGAACAAATC	GATTGGAAA	GTAAAATTAA	TTTCTATAAA	TGTTTGTGCA	9240	
ATTGTTTCG	GT ACTATTTAG	ATTCA	GTCTA	CTATATACAA	TATTTCCGGA	ACATTCAACT	9300

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TTTTAACTCT ATTTATTACT AGATTCATA ATTAAAAAAC CTACTGACCA AGCTAGAAAG	9360
CTTGATACAA TAGGCTTTT AAAGACTGAT TATTTAACAG CGTCTTAAG AGCTTTACCA	9420
GCTTGAATG CTGGTACTTT AGAAGCTGCA ATTGTCAATT CTTTACCAAGT TTGTGGTTG	9480
CGACCTTAC GTTCTGCGCG CTCACGAAC TCAAAGTTAC CAAAACCGAT CAATTGAAC	9540
T	9541

(2) INFORMATION FOR SEQ ID NO: 133:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 3502 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 133:

TTGACTATCC TATCATGCTT TCTAAGGTCT ACTCAAGAAA ATCATTTCAC	60
CCTTTCTCAA AAAAGTTAAA AAATTTCTC AAAAACGCTT GACTCTGACC TAAGGCGAAG	120
GGTTATACTA TCATTGTAAG GAGGAAATCA TGTACCATAT AAAAGAACGT GCGCAGCTTT	180
CAGGTGTCTC TGTCAAGACC CTGCATCACT ATGACAAGAT AGGACTCTTG GTCCCCTTAA	240
AGTCGGAAA CGGCTATCGA ACCTACAGTC AAGAGGATTG GGAACGCCTT CAGGTCAATT	300
TTTACTACAA ATATCTAGGC TTTCTTTAG AGAAAATAGC AGAGCTGTTA AAGGAAGAAA	360
GGACAGATT ATTGCCCAT TTGACTAGGC AGTTGGACTA TCTAACTCGC GAAAGGCAAC	420
ATCTGGATAC CTTGATTTCC ACCTTGCAAA AACTATTCA AGAACAAAAA GGAGAAAGAA	480
AAATGACCAT TGAGGAAAAA TTCACGGGAT TTAGCTATCA AGACAATCAA AAATACCACC	540
AAGAAGCGGT AGAGAAATAT GGTCAAGAAG TCATGGGACA AGCGCTCGA CGCCAAAAAG	600
GTCACGAAGA CGAGGCTACG GCCGCCTTA ACCAAGTCTT TCAAACTTG GCACAAAATC	660
TTCAAGTTGG TTTACCTGCA ACAGCAACCG AAAACCAGGA GCAAGCAGCC AAGCTCTTGC	720
AAGCCATTGCA CACTTATGGA TTGACTGCT CTATTGAGGT ATTGGTCAAT ATCGTAAAG	780
GTTACGTCTA CAACCCAGAG TTAAAGGAAA ACATTGACAA GTTGGTTCT GAAACAGCCC	840
AGTACACGTC AGATGCCATT GCGGTTACG TTCAGACAAA TGCAGAATAA ATAGGCTAGG	900
AATTCCTAG CCTATTTTT ACTTCAAATC ATAAAGCCAG TCGTCACCGT TTTTAGTA	960
AAAGAATTCA CTGAGATCTT CTTCTAGAAA CACACGAAGC ATATCAGACA TATCATCGGT	1020
TGCAAGTTT AGATGAGAAA GATTTTCAAA GTCCTCCCAC CAAACTTCC CTTCGTCTGA	1080
AGACTGGAGT TCACCAAGTAA AGTGTCTGT CTTGTAAAAA AGGACGACAT AACGATAATC	1140

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CTTGTGTC	TACCA	GTTTG	GGGT	TTGGAA	ATGA	TCAGACC	AGT	1200									
TTCTTCTT	TC	ACTTCACGAA	TGACAGC	ATC	GACAAAGG	AT	TCGCCAC	TT	CAACATGACC	1260							
ACCAGGAAA	AA	GTAATGCCAG	ACCAGTC	GGG	ATTA	ACTCGG	TCTTGGACCA	GGAC	CTTATC	1320							
TCCGTTT	TA	ATCATACACA	TGTTAACAAA	TTCGACTGCC	TCTCTTCTGT	TCATTCTTCA	1380										
CAACCTT	AA	TCTTAATCA	TAATGCAGAC	TTCCC	CCAC	CCAGCCG	TA	CAGAGGGCAG	1440								
AAGTGA	TGTT	AAAGCCACCC	GTG	GGGCAT	TGATATCC	AACTTC	CGCCT	GCAAAGTGG	1500								
GGCCAGGT	AC	CAGCTTACTT	TCA	AGGGTTT	TAGGATTGAT	TTC	CTTGAGA	CTGACTCCAC	1560								
CCTTGGT	AA	AAAGGACTTT	GCA	AGGGACA	TTTTCC	AGT	TACAGGA	ATT	TTAAGTTCTT	1620							
TAATGG	AC	TG	GACAAGTTGT	TCTCG	TTCTC	CT	TTTCAG	TGACT	TTTCAGGAT	1680							
ATCCTT	GT	TAC	AAAAAATT	CG	GCC	AAAGCG	TT	CTG	TAACAA	GGTTTT	AAA	GC	GT	TTTCA	1740		
AGGATTT	TTC	CCG	ATTT	TCT	AGAA	ATG	TAACCA	AGTC	CTT	CTCAG	AA	AG	TGAGG	CA	1800		
AAACATCG	AG	TG	AGAGA	ACC	TCCC	CAC	TT	TGAC	AAAG	GCT	AGAC	ATGCG	TAGG	CAGCAG	1860		
GACCTG	AC	AA	ACCA	AGTGG	GTAA	AGAGT	AA	TCA	GTGAG	ATG	GATG	ACATG	ATGC	TTACCA	1920		
TTAGGG	TC	AC	ATCG	TCCAGA	GAA	ATAC	CTT	GTA	AGG	CTT	ATG	GGAAA	TCT	GTTA	ATA	1980	
AAGGACT	TT	TC	AGCAGC	CTCA	AG	ATCGG	TGA	TGG	TATG	C	TT	AAA	ATGG	CGA	GCA	ATCTCG	2040
GACCA	AA	AC	AGTC	GAAC	GTC	GAAGG	AT	AGACT	TAC	AC	CTG	TG	ACA	ATGAG	TT	2100	
TCTCACA	AG	GT	AAAG	TTG	TG	CGT	GACT	TA	AGG	AC	AA	CTG	TC	CAT	CT	2160	
CAGAAACG	AT	TT	CT	CT	GT	TTG	GATT	TT	CGG	GT	TT	CTT	CCA	TT	CTT	2220	
AAGCTT	CG	AT	AA	TAGT	CCG	A	GACT	GTC	AC	TG	TT	GG	CGA	AA	TCG	2280	
CCTTAAG	TT	AA	AC	AC	CC	TT	TCT	GAA	AG	CTG	AT	CT	GG	CA	ATG	GGG	2340
AGAAA	AC	CT	GTAA	AGAA	CGT	CCG	TT	CAG	ATT	CC	AG	CTA	GG	AG	TCA	AGC	2400
TACCA	TT	GT	GG	TG	TC	AC	TTG	TC	GG	CT	TT	CC	AA	GG	TT	CT	2460
TCCGAT	TTT	TT	TT	TC	CG	AT	GAG	AGG	TTT	CT	TT	CC	ATA	AAA	G	ATCGTAGCCA	2520
TCATACCAG	C	AG	GT	CCCC	CC	CG	ATG	ACAA	TAG	TAT	CAA	ATG	TTT	CATA	G	CTCTATTGT	2580
ACCAC	AA	AA	AA	ACA	AG	AG	AT	GTC	AC	T	CTG	CA	AG	AA	TT	CA	2640
TAGCC	CC	AT	CA	GC	AA	AC	CG	CC	T	C	TCT	CTG	CA	TA	AC	AG	2700
ATTTA	AT	AT	AT	CCG	CT	GT	GG	GA	AG	GT	TT	CG	AT	GAG	AC	AT	2760
TTTCGTT	AT	AT	AT	TG	CG	TT	GG	GA	AG	TT	CA	GG	AC	CC	AG	AC	2820
TCATCATAG	GG	AG	CAG	CTT	GA	AC	TG	AT	TT	CTT	G	TG	CC	TT	G	TA	2880

908	
GTCCCAGTTT CACTAGCTTT TCCGACCATA CGAATGTTGA GAAGGCCAAC GACCGTACCG	2940
ATAAGCTTGC TCAAACGGCC GTTCTTCACC AAGTTATCGA CTGGCTAG GACAAGAGC	3000
AACTTAGTTT TTTCTTGATA GGCGGTGATA GCTTCAACCA CTTCTTCAAA AGACAAGCCC	3060
TGGTCAATCA AGTCATTCAA TTTTCTACG AGTAGGTCAA CTTCACCCACC AGCAGATAAA	3120
CTATCAATCA CATGAATCTT AGTGTCAAGGA TGGTCTTCCA GATAAAATATT CTTGCTAGT	3180
TGAGCACTAT TGTGACTGCC AGAAAGGGTA CCTGTGATGG TTACTAGGAA AATGTTTTG	3240
GCACCTTCAA ATGCTCGCAA ATAGTCATCT GGGCTTGGAC AAGCCGATTT TGAAGCTTCT	3300
GCAGTTGCAT ACATGGTTTC CATCATTGG TCAATATCGA GACTGGCGTC ATCAACAAAG	3360
ACCTGATCAG CTACTTGAAAT GGTAAAGGGG ACACTTACAA AGGTTGTGTT AATAGCTGGT	3420
GTTGGCAGTT GACGATAATC ACAACCAGAG TCAGCAATAA TCTTCCAAGT CATAGAAATT	3480
CTCCATCTTT GTCAGGAACG AT	3502

(2) INFORMATION FOR SEQ ID NO: 134:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 12665 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 134:

CGATTGATT TTTTAAAGCG TTCGATAGAG AATGAGAAC GAATCCTTAG CAATGGCGGG	60
AAAGAATTG GAGTTGAGAA TACAAAACGA TTAACCTATGG CTCATATTGT TTTTTATCTC	120
TCTTGCTTGG TTGAGGCAAT GGTGACAAG ACAATTTTG ATGGCATGGG CATGGTTGGT	180
TTAGTCTTGC TTATTTTTTC TATGCTGATG TTGATGTTGG TGATTCACTT GTTGGGAGAT	240
ATTTGGACAG TGAAGCTTAT GCTTGTCAAT AACACAAAT ATGTAGATCA TATCTGTTT	300
AGGACAGTAA AACACCCCAA TTACTTTTA AATATTCTTC CTGAGTTGAT TGGCTTGACC	360
TTGTTGAGTC ATGCTTATGT GACTTTGTT TAGTTTTTC CAGTTATGC AGTTATTTG	420
TATCGACGAA TAGCTGAAGA GGAAAGCTA TTACATGAAG TTATAATCCC AAATGGAAGC	480
ATAAAGAGAT AAATACAAAA TTCGATTTAT ATACAGTTCA TATTGAAGTG ATATAGTAAG	540
GTTAAAGAAA AAATATAGAA GGAAATAAAC ATGTTGCAT CAAAAAGCGA AAGAAAAGTA	600
CATTATTCAA TTCGAAATT TAGTGTGGA GTAGCTAGTG TAGTTGTTGC CAGTCTTGT	660
ATGGGAAGTG TGGTTCATGC GACAGAGAAC GAGGGAGCTA CCCAAGTACC CACTTCTTCT	720
AATAGGGCAA ATGAAAGTCA GGCAGAACAA GGAGAACAC CTAAAAAAACT CGATTCAAGAA	780

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CGAGATAAGG	CAAGGAAAGA	GGTCGAGGAA	TATGTA	AAAAA	AAATAGTGGG	TGAGAGCTAT	840
GCAAAATCAA	CTAAAAGCG	ACATACAATT	ACTGTAGCTC	TAGTTAACGA	GTTGAACAAAC		900
ATTAAGAACG	AGTATTGAA	TAAAATAGTT	GAATCAACCT	CAGAAAGCCA	ACTACAGATA		960
CTGATGATGG	AGAGTCGATC	AAAAGTAGAT	GAAGCTGTGT	CTAAGTTGA	AAAGGACTCA		1020
TCTTCTTCGT	CAAGTTCAGA	CTCTTCCACT	AAACCGGAAG	CTTCAGATAC	AGCGAAGCCA		1080
AACAAGCCG	CAGAACCAAGG	AGAAAAGGTA	GCAGAAGCTA	AGAAGAAGGT	TGAAGAAGCT		1140
GAGAAAAAAAG	CCAAGGATCA	AAAAGAAGAA	GATCGTCGTA	ACTACCCAAC	CATTACTTAC		1200
AAAACGCTTG	AACTTGAAAT	TGCTGAGTCC	GATGTGGAAG	TTAAAAAAGC	GGAGCTTGAA		1260
CTAGTAAAAG	TGAAAGCTAA	CGAACCTCGA	GACGAGCAAA	AAATTAAGCA	AGCAGAAGCG		1320
GAAGTTGAGA	GTAAACAAGC	TGAGGCTACA	AGGTTAAAAA	AAATCAAGAC	AGATCGTGAA		1380
GAAGCAGAAG	AAGAAGCTAA	ACGAAGAGCA	GATGCTAAAG	AGCAAGGTAA	ACCAAAGGGG		1440
CGGGCAAAAC	GAGGAGTTCC	TGGAGAGCTA	GCAACACCTG	ATAAAAAAGA	AAATGATGCG		1500
AAGTCTTCAG	ATTCTAGCGT	AGGTGAAGAA	ACTCTTCCAA	GCCCATCCCT	GAAACCAGAA		1560
AAAAAGGTAG	CAGAAGCTGA	GAAGAAGGTT	GAAGAAGCTA	AGAAAAAAGC	CGAGGATCAA		1620
AAAGAAGAAG	ATCGCCGTAA	CTACCCAACC	AATACTTACA	AAACGTTGA	ACTTGAAATT		1680
GCTGACTCCG	ATGTGGAAGT	TAAAAAAGCG	GAGCTTGAAC	TAGTAAAAGA	GGAACGCTAAG		1740
GAACCTCGAA	ACGAGGAAAA	AGTTAAGCAA	GCAAAAGCGG	AAGTTGAGAG	TAAAAAAGCT		1800
GAGGCTACAA	GGTTAGAAAA	AATCAAGACA	GATCGTAAAA	AAGCAGAAGA	AGAACGCTAAA		1860
CGAAAAGCAG	CAGAAGAAGA	TAAAGTTAAA	GAAAAACCAG	CTGAACAAAC	ACAACCAGCG		1920
CCGGCTCCAA	AAGCAGAAAA	ACCAGCTCCA	GCTCCAAAAC	CAGAGAATCC	AGCTGAACAA		1980
CCAAAAGCAG	AAAAACCAGC	TGATCAACAA	GCTGAAGAAG	ACTATGCTCG	TAGATCAGAA		2040
GAAGAATATA	ATCGCTTGAC	TCAACAGCAA	CCGCCAAAAAA	CTGAAAACC	AGCACAACCA		2100
TCTACTCCAA	AAACAGGCTG	GAAACAAGAA	AACGGTATGT	GGTACTTCTA	CAATACTGAT		2160
GGTTCAATGG	CGACAGGATG	GCTCCAAAAC	AATGGCTCAT	GGTACTACCT	CAACAGCAAT		2220
GGCGCTATGG	CGACAGGATG	GCTCCAAAAC	AATGGTTCAT	GGTACTATCT	AAACGCTAAT		2280
GGTTCAATGG	CAACAGGATG	GCTCCAAAAC	AATGGTTCAT	GGTACTACCT	AAACGCTAAT		2340
GGTTCAATGG	CGACAGGATG	GCTCCAATAC	AATGGCTCAT	GGTACTACCT	AAACGCTAAT		2400
GGTTCAATGG	CGACAGGATG	GCTCCAATAC	AATGGCTCAT	GGTACTACCT	AAACGCTAAT		2460
GGTGATATGG	CGACAGGTTG	GGTGAAGAGAT	GGAGATAACCT	GGTACTATCT	TGAAGCATTCA		2520

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GGTGCTATGA AAGCAAGCCA ATGGTTCAAA GTATCAGATA AATGGTACTA TGTCAATGGC	2580
TCAGGTGCCCT TGCGACTCAA CACAACGTGAT GATGGCTATG GAGTCAATGC CAATGGTGA	2640
TGGGTAAACT AAACCTAATA TAACTAGTTA ATACTGACTT CCTGTAAGAA CTCTTAAAG	2700
TATTCCTAC AAATACCATA TCCTTTCAGT AGATAATATA CCCTGTAGG AAGTTTAGAT	2760
TAAAAAAATAA CTCTGTAATC TCTAGCCGGA TTTATAGCGC TAGAGACTAC GGAGTTTTT	2820
TGATGAGGAA AGAATGGCGG CATTCAAGAG GCTCTTAAG AGAGTTACGG GTTTAAACT	2880
ATTAAGCCTT CTCCAATTGC AAGAGGGTTT CAATCTCTGC CAGGGTGCTG GCTTGCAGAA	2940
TGGCTCCACG GAGTTGGCA GCGCCAGATG TTCCACGGAG ATAGTGAGGA GCGAGACCGC	3000
GGAATTCAACG AACTGCGACG TTTTCTCCTT TGAGGTTAAT CAATCGTTTC AAGTGTTCGT	3060
AGGCGATCTT CATCTTGCTC TCAAAGGTCA AATCAGGTAG GATTTCTCCT GTTTCAAAGT	3120
AATGGTTGAT TTGGTTGAAG AGGTAAGGAT TTCCCATGGC AGCTCGGCCA ATCATGACTG	3180
CGTCAGCACCA AACTTCTTCG ATGCCTTGCT TGGCTTCTTG GACAGTACGG ATATCACCGT	3240
TGGCGATGAA TGGAATCTTG GTTAGAGCTT GGGCAACCTT GTAAAGGGTC TCAAGGTCTG	3300
CGTGGCCAGT ATACATTGAT TCACGGGTAC GCCCATGCAT GCGGAGGGCA GAAACACCTG	3360
CAGCTTCAGC AGCGAGAGCA TTTTCTACTG CAAGAGATGG GTCCGCCAG CCGTACGCA	3420
TTTGACAGT AAGTGGGATA TCAAGGACAG ACTGGACCTT GTTGATGATG GAGTAAATCT	3480
TGTCTGGATC CTTGACCCAC ATAGCACCAG CTTCGTTCTT CACGATTTTG TTGACAGGGC	3540
AGCCCATGTT GATATCGACG ATATCGGTCT TGGTGTTC TTGGATGAAT TCTGCTGC	3600
GTGCTAGGCT GTCTTCATCG CTACCAAAA GTTGGATAGA GACAGGGTTT TCGCCCTCAT	3660
CGATATGAAG CATGTGCAGG GTTTTTCTGT TGTTGTATTG GATTCCCTTG TCAGAGACCA	3720
TTTCCATTAC AACGAGTCCA GCTCCGAGCT CCTTGCGAT AGTACGAAAG GCTGAGTTGG	3780
TCACGCCAGC CATAGGCAGCT AAAACGGTAC GATTGGGAAT CTCATATTG CCAATCATAA	3840
AAGGTGTATT AAGATTTGTC ACGAATGAGT TCCTCCAGGT CCTTTCATC AAAGTTGTAA	3900
GTAGTTGGC AGAATTGACA AGTGATTCTC GCCCCGTGGT CTTCCCTCTT CATTCCCTGT	3960
AAAGTCTGAGC TTGGAAGGCT GGCAAGAGCG TTCATAAAGC GTTCATGGCT ACAGTCACAT	4020
TGGAAACGGA TTTCTCTTC AGAAAGACGC TTGTAGGCTT CGTCCCCGTA GATAGCCTTG	4080
AGGAGGGCTT CGATATGGTC GTGCGTTTCG AGAAGAGTAG AGATAGCTGG CATTCTTGG	4140
ATGCGTTTTT CAAAGCGAGC AATCTCTTCT TTCTGGCTC CTGGCAAGAC TTGAACCTAGG	4200
AAACCACCTG CAACCTTGAC CTTGTCTTCC TCGTCCAAAA GGACATTGAG GCCGACCGCT	4260
GAAGGCAGTTT GTTGGCTTTC AGTAAGGTAA AAGGCAAGGT CTTCACCGAT TTCTCCAGAG	4320

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ATGAGGGGAG TTATAGAGTT GTAAGGATTT CCAGTACCGT AGTCTGTGAT AACGAGGAAT	4380
TGACCATTTC CAACAAAAGG TCCGACTAGG ACTTCACCAG TCGCAGTCTT TTTGATGTCA	4440
ACACCAGGAT TTTGAACATA GCCTTGACG TTCCCTTGG TATCAGCGAC GGTGATAATA	4500
GCACCTAGAG AGCTAGATCC CAACACCTTA ACTGTAAGTT TGGTATTTCC TTTTCATTG	4560
GCTGCGAGAA TCTGGCTAGC GATAAGAGTT CGACCCAAGCG CTACAGTTGA GCTAGCTTGG	4620
GTTTGATGTT TTTCTTGAGC AGTGCAGGACG GTTTCAGTGC TATCAAGGAC AAAAGCACGA	4680
AAGGcTCCGC TTTCTGATAT AGTTTAATA ATTTTATCCA TAGCTACTAT TTTAGCATAA	4740
AAATGCCCAA AGGGGGAGCC GTGTGTTTAC TGATTTCAG GATAATGGAC CAGGAAATCA	4800
GCATGAAAAT AAAAAGAGAA ACAGATTATT TTAGCATTG TCAGATTAT GCTATGCTTA	4860
AGGTAGAAAA TGAAAGGGAT AACAAATGTA TTTAGGAGAT TTGATGGAGA AAGCCGAGTG	4920
TGGTCAATT TCAAATACCTT CCTTCTATT ACAAGAGTCT CAGACGACCG TCAAGGCTGT	4980
AATGGAAGAA ACAGGATTTT CAAAAGCAAC CCTAACCAAA TATGTCACCC TGCTCAATGA	5040
CAAGGCTTTG GATAGTGGCT TAGAGCTGGC TATTCACTCA GAAGATGAAA ATCTGCGTCT	5100
GTCTATCGGT GCAGCTACCA AGGGGAGAGA TATTGGGAGC TTGTTTTGG AGAGTGCTGT	5160
TAAATACCAAG ATTTTGGTTT ATCTTCTCTA CCACCAACAG TTTTTAGCCC ATCAGCTGGC	5220
TCAAGAATTG GTGATTAGCG AGGCTACGCT TGGTCGTAC TTGGCTGGTT TAAATCAGAT	5280
TTTGTCAAGAA TTTGATTAT CCATCCAAA TGCCGTTGG CGAGGTCCAG AGCATCAGAT	5340
TCACTATTTC TATTCTGTC TTTCCGAAA GGTCTGGTCG AGTCAGGAAT GGGAAAGGTCA	5400
CATGCAGAAA CCAGAGAGAA AACAGGAGAT TGCCAATTAA GAGGAATCT GCGGTGCAAG	5460
TTTGTCTGCG GGGCAGAAAT TGGACTTGTT TCTCTGGCT CACATCAGTC AACAAACGTCT	5520
TCGGGTCAAT GCTTGTCACT TTCAGTCAT AGAAGAGAAA ATGCGAGGGT ATTTTGACAA	5580
TATCTTTAT CTTCGTTGC TGAGAAAGGT TCCGTCCTTT TTTGCTGGGC AACATATTCC	5640
ACTAGGAGTT GAGGATGGTG AGATGATGAT ATTCTTCTCT TTTCTCCTAT CTCATCGCAT	5700
TCTTCCTCTT CATACTATGG AGTATATTCT TGGTTTGGA GGGCAGTTGG CAGATTACT	5760
GACGCAATTG ATTCAAGAAA TGAAGAAGGA GGAACATTG GGGGATTATA CAGAGGACCA	5820
TGTCACCTAT GAACTCAGTC AGCTTGTGC TCAAGTCTAT CTCTATAAGG GCTATATTTC	5880
ACAGGATCGC TACAAGTACC AGTTAGAGAA TCGTCATCCA TATTACTGA TGGAACATGA	5940
TTTTAAAGAG ACAGCAGAGG AGATTTTCA TGCTCTACCT GCTTTCAAC AGGGGACAGA	6000
TTTAGATAAG AAGATTCTCT GGGAAATGGCT CCAGTTAAC GAATATATGG CTGAAAACGG	6060

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TGGCCAGCAT	ATGCGGATTG	GTCTGGATTT	GACATCTGGT	TTTCTTGCT	TTTCAAGGAT	6120
GGCAGCCATT	TTGAAACGGT	ATTTGGAATA	CAATCGTTT	ATTACCATTG	AAGCTTATGA	6180
CCCTAGTCGG	CATTATGATT	TGCTGGTTAC	CAATAACCCG	ATTCTAAGA	AGGAACAGAC	6240
ACCAGTCTAT	TATTTAAAAA	ATGACTTGGA	TATGGAGGAT	TTGGTAGCGA	TTGCCAGTT	6300
ATTATTCACT	TAAAAGGCTT	GGTTAATCCA	GGTCTTTTT	GTGAAATTCA	CACAATCTCC	6360
TCACATTTTT	TTAAAAATTA	AAAAAAAGTTG	ATAAACAAAGA	AAGCGCTTTA	TTTTGTATAC	6420
TAGTAAGTGT	AAAGAGGAAA	CACCTCAAGA	TCTTTATCAG	GAGGACAGTA	CATGTCACAA	6480
GAAAAATACA	TCATGGCCAT	TGACCAGGGA	ACTACAAGTT	CTCGTGCCAT	CATTTCAAC	6540
AAAAAAGGGG	AAAAGGTAG	CTCGAGTCAA	AAAGAGTTA	CCCAGATT	CCCTCAGGCA	6600
GGTTGGGTTG	AGCACAATGC	CAATGAAATT	TGGAACTCTG	TTCAGTCAGT	TATTGCGGGT	6660
GCTTCATCG	AAAGTGGTGT	CAAGCCAAAT	CAAATCGAGG	CAATCGGGAT	TACCAACCAA	6720
CGTGAAACAA	CGGTTGTCTG	GGATAAGAAA	ACAGGACTTC	CTATCTACAA	TGCTATCGTT	6780
TGGCAGTCAC	GCCAGACAGC	ACCTTTGGCT	GAGCAACTAA	AAAGCCAAGG	TTATGTGGAA	6840
AAATTCCATG	AAAAGACTGG	TTTGATTATT	GATGCTTACT	TCTCTGCTAC	CAAGGTCGTT	6900
TGGATTTGG	ATCATGTAGA	AGGTGCTCAA	GAGCGAGCAG	AAAAAGGGGA	ATTGCTCTTT	6960
GGTACTATCG	ATACTTGGTT	GGTTTGAAA	TTGACTGACG	GTGCGGCTCA	CGTGACTGAC	7020
TACTCAAATG	CAGCTCGTAC	CATGTTTAT	AACATTAAG	AACTCAAATG	GGATGATGAG	7080
ATTTTGGAAA	TCCTTAACAT	TCCGAAGGCT	ATACTTCCAG	AAGTCGTTTC	TAACTCCGAA	7140
ATCTACGGCA	AGACAGCTCC	ATCCATTTC	TACGGTGGAG	AGGTGCCAAT	CTCAGGTATG	7200
GCTGGGGACC	AAACAAGCAGC	CCTCTTGGGA	CAGTTGGCTT	TTGAGGCCAGG	TATGGTTAAG	7260
AATACTTATG	GAACAGGCTC	TTTCATCATC	ATGAATACTG	GGGAAGAGAT	GCAGTTGTCT	7320
GAAAACAACC	TCTTGACAAC	CATTGGTTAC	GGAATCAACG	GTAAGGTTA	TTATGCCTTG	7380
GAAGGTTCTA	TCTTCATCGC	AGGAAGTGCT	ATTCAGTGGC	TTCGTGACGG	TCTTCGCGATG	7440
GTTGAAAATT	CACCAGAACAT	TGAAAAATAC	GCTCGTGATT	CTCACAAACAA	CGATGAAGTT	7500
TATGTCGTT	CAGCCTTAC	AGGTCTAGGC	GCTCCATACT	GGAACCAAAA	TGCTCGTGGT	7560
TCCGTCTTG	GTGGACTCG	TGGAACAAGC	AAAGAAGACT	TTATCAAGGC	GACTTTGCAA	7620
TCTATTGCTT	ATCAAGTGC	TGATATCATC	GACACCATGC	AAAGTGGATAAC	TCAGACCGCC	7680
ATTCAAGTAC	TGAAGGTGGA	TGGTGGTGCA	GCCATGAACA	ACTTCCTCAT	GCAGTCCAG	7740
GCGGATATTT	TAGGCATGTA	CATTGACACGT	GCTAAAAACC	TGGAAACAAC	AGCTCTAGGA	7800
CGGGCCTTCC	TAGCAGGTTT	GTCAGTAGGG	TACTGGAAAG	ACTTGGACGA	GTTGAAACTC	7860

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TTGAACGAGA CAGGAGAACT CTTTGAGCCA TCTATGAACG AATCTCGCAA GGAACAACTC	7920
TACAAGGGCT GGAAGAAGGC TGTGAAAGCA ACTCAAGTCT TTGCGGAAGT AGACGACTAA	7980
TACTGGCAGA ATAAAGCGAT TTATTAGAA AGTGTGTAAA TATGGAATT TCAGGAAAAA	8040
CACGTGAATT GTCAATTAAA AAAATGCAGG AACGTACCCCT GGACCTCTTG ATTATCGGTG	8100
GAGGAATCAC AGGAGCTGGT CTAGCCTTGC AGGCCGCAGC TAGCGGTCTT GAGACTGGTT	8160
TGATTGAAAT GCAAGACTTT GCAGAAGGAA CATCTAGTCG TTCAACAAAAA TTGGTTCACG	8220
GAGGAACCTCG TTACCTCAAA CAATTGACG TAGAAGTGGT CTCAGATACG GTTCTGAAC	8280
GTGCAGTGGT TCAACAAATC GCTCCACACA TTCCAAAATC AGATCCAATG CTCTTACCA	8340
TTTACGATGA AGATGGAGCA ACCTTAGCC TCTTCCGTCT TAAAGTAGCC ATGGACTTGT	8400
ACGACCTCTT GGCAGGTGTT AGCAACACAC CAGCTGCGAA CAAGGTTTG AGCAAGGATC	8460
AAAGTCTTGGA ACGCCAGCCA AACTTGAAGA AGGAAGGCTT GGTAGGAGGT GGAGTGTATC	8520
TTGACTCCG TAACAACGAT GCGCGTCTCG TGATTGAAAA CATCAAACGT GCCAACCAAG	8580
ACGGTGCCTT CATTGCCAAC CACGTGAAGG CAGAAGGCTT CCTCTTGAC GAAAGTGGCA	8640
AGATTACAGG TGTTGTAGCT CGTGATCTCT TGACAGACCA AGTGTGAA ATCAAGGCC	8700
GTCTGGTTAT TAATACAACA GGTCCCTTGGA GTGATAAAAGT ACGTAATTG TCTAATAAGG	8760
GAACGCAATT CTCACAAATG CGCCCAACTA AGGGAGTTCA CTTGGTAGTA GATTCAAGCA	8820
AAATCAAGGT TTCACAGCCA GTTTACTTCG ACACAGGTTT GGGTGACGGT CGTATGGTCT	8880
TTGTTCTCCC ACGTGAAAAC AAGACTTACT TTGGTACAAC TGATACAGAC TACACAGGTG	8940
ATTTGGAGCA TCCAAAAGTA ACTCAAGAAG ATGTAGATTA TCTACTTGGC ATTGTCAACA	9000
ACCGCTTCCC AGAATCCAAC ATCACCATTG ATGATATCGA AAGCAGCTGG GCAGGTCTTC	9060
GTCCATTGAT TGCAGGGAAC AGTGCTCTG ACTATAATGG TGAAATAAC GGTACCATCA	9120
GTGATGAAAG CTTTGACAAC TTGATTGCGA CTGTTGAATC TTATCTCTCC AAAGAAAAAA	9180
CACGTGAAGA TGTTGAGTCT GCTGTCAGCA AGCTTGAAAG TAGCACATCT GAGAACATT	9240
TGGATCCATC TGCAGTTCT CGTGGGTCTA GCTTGGACCG TGATGACAAT GGTCTCTGA	9300
CTCTTGCTGG TGGTAAAATC ACAGACTACC GTAAGATGGC TGAAGGAGCT ATGGAGCGCG	9360
TGGTTGACAT CCTCAAAGCA GAATTGACC GTAGCTTTAA ATTGATCAAT TCTAAAAC	9420
ACCCCTGTTTC AGGTGGAGAA TTGAACCCAG CAAATGTGGA TTCAGAAATC GAAGCCTTG	9480
CGCAACTTGG AGTATCACGT GGTTGGATA GCAAGGAAGC TCACTATCTG GCAAATCTTT	9540
ACGGTTCAAA TGCACCGAAA GTCTTGCAC TTGCTCACAG CTTGGAACAA GCGCCAGGAC	9600

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TCAGCTTGGC AGATACTTTG TCCCTTCACT ATGCAATGCG CAATGAGTTG ACTCTTAGCC	9660
CAGTTGACTT CCTTCTTCGT CGTACCAATC ACATGCTCTT TATGCGTGAT AGCTTGGATA	9720
GTATCGTTGA GCCAATTGG GATGAAATGG GACGATTCTA TGACTGGACA GAAGAAGAAA	9780
AAGCAACTTA CCGTGCAGAT GTGAAAGCAG CTCTCGCTAA CAACGATTAA GCAGAATTAA	9840
AAAATTAAGA AAAAATAAAA GAGGTGGAGG GCAGCATTCC TTGTCGCCCG TCCCTCTTT	9900
TTAATGGAGA CAGAAAGATG ATGAATGAAT TATTTGGAGA ATTTCTAGGG ACTTTAATCC	9960
TGATTCTTCT AGGAAATGGT GTTGTGCAG GTGTGGTTCT TCCTAAAACC AAGAGCAATA	10020
GCTCAGGTTG GATTGTGATT ACTATGGGTT GGGGGATTGC AGTTGCGGTT GCAGTCTTG	10080
TATCTGGCAA GCTCAGTCCA GCTTATTTAA ACCCAGCTGT GACCACGGT GTGGCCTTAA	10140
AAGGTGGTT GCCTTGGGCT TCCGTTTGC CTTATATCTT AGCCCAGTTC GCAGGGGCCA	10200
TGCTGGGTCA GATTTGGTT TGGTTGCAAT TCAAACCTCA CTATGAGGCA GAAGAAAATG	10260
CAGGCAATAT CCTGGCAACC TTCAGTACTG GACCAGCCAT CAAGGATACT GTATCAAAC	10320
TGATTAGCGA AATCCTTGGAA ACTTTGTTT TGGTGTGAC AATCTTGCT TTGGGTCTTT	10380
ACGACTTTCA GGCAGGTATC GGAACCTTGT CAGTGGGAAC TTTGATTGTC GGTATCGTC	10440
TATCACTAGG TGGGACAACA GGTTATGCCT TGAACCCAGC TCGTGACCTT GGACCTCGTA	10500
TCATGCACAG CATCTTGCCA ATTCCAAACA AGGGAGACGG AGACTGGTCT TACGCTTGGA	10560
TTCCCTGTTGT AGGCCCTGTT ATCGGAGCAG CCTTGGCAGT GCTTGTATTC TCACTTTCT	10620
AGTTTATACT CTTCGAAAAT CAAATTCAAAC CCACGTCAGC GTCGCCTTAC CGTACTCAAG	10680
TACAGCTTGC GGCTAGCTTC CTAGTTGCT CTTGATTCTT CATTGAGTAT TAGAAAACAA	10740
TTATGTTGAT AGAGCTTGGG CAAGAGCCA ATTTCAGCAA AAAATGAAGT AAATCTCTC	10800
ATAATAAAAC GCATCATATC AAGCACGAAA ATTCCACGAG GTCAACTACA GTCAGAAAGC	10860
TGAACAAACAA GCCAAAACGC CCAAAAAAGG CGGAAAAAG CAAGCACCTG CAAGCAACGT	10920
GCCGAAATGG TCAAATCCTG ATTATGTCAA CGAATTAGAC CCAAAATCG TTGATATGCT	10980
AGTAGAATTTC CACAAGTCAC AAGGCACTTT GGAAACTCCC GAGGCAGCAAG CAGAAATCG	11040
CCAAAAACGT GAAGAAATCG AGCAAAGGAG AGCTGAGCTT GAGGGTAAAA ACAAGAGCT	11100
TTTGAACCGC TTGAACAAAT AGAGTTCGC AAGTATTATG CTTACAAATT ACTTGAGCAA	11160
TTAACTAAAA TATAAACCT GCCTTATAT CTAGGCAGGG TTTATATTTT AGAAATTAC	11220
GTAGGTTGTT ACGGTTTTA CATAACCCAGT ATAGTTGAG TTTCTATAGT ATTCACTGAT	11280
AAACTTCCAT TTTCTTGAG CAACATGGAT ATAAGTACTT GTTATGAGT ATGGATATGG	11340
GCTTGTGAA TCCAAGTAAG ACTGATAAGC TTGTATACCA AAATATGCTC CACCAATTAT	11400

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TGCACCCCAT	GGACCCCCA	ATAAACGACC	TATCCTACCA	ATCATATAAC	TGATTCCAGC	11460
ACCAGTCATG	AAGTTAGCGA	ATGTGTTAGC	TTGTTTATTC	CCATGTATTG	TGTTGACGTA	11520
ATTCCAAACA	TTAGGATCGT	ATGATCTAAA	AGATATATTT	AGGTCGATTT	CATTCTTTG	11580
ATAAGCCATA	TAAAATGCC	CATTGATATA	GACGCCGTCA	GCACGTCGTT	CAATAGTGTC	11640
TACACTTCCA	TCTGGATTGA	CAACCTCAAG	AACTTCATCG	CTTAAAATAT	TTACTTGCGT	11700
ATCTCCGAAAC	CGCACTGATG	AGCCATTCTC	AAACTGAGCC	TCACCAGATA	CAACTTTAGA	11760
GTGTTGCCGAT	AAGCTATCAT	CAGCAAAAAC	AAACAAGCGA	CGGGGAAATG	CTAGACATAC	11820
AGAAAACAGA	CATAACTAGC	AAACACATGC	ATTTAAACAT	CTTAGACATA	ACGGAAACTC	11880
CTTTGTATTT	TTGATTTTT	TCAACTTTA	TTATACAATA	AAACCAAATA	AAAAGAAAGC	11940
GGTAACAATA	TGCTTAATGC	GAAAATTTT	TATATATTTT	TATGTTTGAT	CGTTATCGAA	12000
ACTACAGGCT	TGTTGTGTT	GAAAAGAGGT	CTCGAAATGG	GTTATTTAGA	CACAGAAGCT	12060
ATTATCCTCG	CAGTTTTTC	ATTTGCTTTT	TACAACCTAT	GTTCATTGCG	TTGGGTCTGC	12120
TCTACAATAA	AAAACAATAA	AAAATAAATA	GACGTATTTT	CAAAAAAAAC	maAATGCATA	12180
TTTATATTAG	CAAACGACG	ATTTAAATCG	TCGTTTTTTT	GTTAGTACGAC	GGGCATGTCG	12240
TATATCTGAG	GTGTAAGTCC	TCAGCCTGAC	TATCGTGAGG	TAGCAGGGAG	AGGAAGGGAT	12300
AGCGAAATCG	TGGCTCTACG	AACAGGAACG	TGATAGTAAG	GCGTATATAG	CGGATAAGGA	12360
GGCTTCAAAAC	TCTAAAGTCC	AAAAGGTAG	TCGTAACCTA	TATGTTGAAA	TCACGAGAGT	12420
AATTGAATTC	GGACTAAGGT	TTGTGTGAAA	AAGATAAAATC	TTTCTAGAGT	CTAAAGACTC	12480
TGCGTCAGAT	TTCCTATTTT	CACTGTAACC	TTTTAACGTC	CTCATATCTT	GTATAAACGA	12540
GGAAAGATGT	ACGACTTATC	CCGTGAGGTT	TCATGAGCGT	GAAAGCGTAG	TAACAACGAA	12600
TCATGAGAAG	TCAGCCGAGC	CCATAGTAGT	GAGGAAACTT	CCGTAATGGA	AGTGGAGCGA	12660
AGGGG						12665

(2) INFORMATION FOR SEQ ID NO: 135:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5305 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 135:

CGCTAACATCAC TACAATCATT TTATGTTACT TTTTCACTCT CAAGAAAAGC AAGAAGTATT

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916

CATTTTAGTT	TCATTTAGTA	TTATTTGCA	TACCTAAAAT	ACAGTAAAAA	ATCAGTCATC	120
TTGGTATGCT	CCTGCTTC	CTATTCAACA	CGTTTTGAC	TTATACTAGG	CTCATTCCA	180
AAAGCATTAT	ATAATAGTGA	TATGAAACCA	ACTAAACTAA	ACAAGAAATA	TAAGCAATAA	240
AAATTCTTT	AAAAGATCTT	ACTAAAGCTA	ATACTAAATA	AAAATAAAAG	AGTAAACTAG	300
GAAGTTTATT	TCAAACAACC	TAAAATACTG	ATTTCGGCT	GAAGATAATA	CTGGAGTGCA	360
AATTAATGGG	GTTATAATAA	ATAGCTGATA	GCTTGTGTTG	GTTTGGATT	TTTTAAGAGT	420
AGATGAGTAT	AAAACATATA	AGGAGGACGA	AGGTGGCTAA	AAATTTAAAA	TTAAAATTAG	480
CTCGGGTAGA	GCGTGATT	ACACAAGGTC	AACTGGCAGA	GGCTGTCGGG	GTGACACGCC	540
AGACTATTGG	TTAATAGAG	GCGGGAAAT	ACAATCCCAG	TCTCTCGCTC	TGCCAGTCTA	600
TTTGCAGATG	TTTAGGGAAA	ACCCTAGACC	AACTATTTG	GGAGGAAGAA	GATGAAAAAT	660
AGATTTTATT	ATTCTCAATT	ACTAGACGAA	AGAGAAGAAC	AACTGTTCAA	TAAAGCGGGC	720
TCTGAAAGTT	TCTATATCTG	CATTGCTTG	TCGCTCCTAT	CTTATATCAT	TTCACTATT	780
GCACCAAGCC	TTTTAATTC	TAATATGCTG	CTAACCGTTA	TCATCATAGG	GACATTTCAC	840
TTTTCAATC	GTGCCCGTTA	TCTGGGAGTG	ACCTACTATG	GTCGTTTCA	TTTACGATT	900
TGGGTTGTT	TTTCCTAAC	CTTGGCTATT	ACGGCTCTTT	TGATGTTGCA	GAATTATCAA	960
TTCAACATAG	AAATTTATCA	GCACAATCCT	TTGAATTTTA	AATACCTGTC	TGCTGGGTC	1020
ATTACTTATA	TCATTTACCT	TCCGTGGATC	TTTATTGGCA	ATCTTGGCT	TAAGAGCTAT	1080
GGCGAATGGG	CTCAGAAAAA	ATTGAACAA	GATATGGATG	AATTGGAGAG	TGGAGAACAG	1140
CTTGGTTACTC	TTTCTCAAT	CCAGCTAAA	TGTGATATAA	TAGTACTAAT	TTATTGGAAT	1200
ACATGAAAGT	TCTTGAAAAT	TTTCATGGGT	TTCTAGCTAA	GGAAGTAGGA	AAAGTATGTA	1260
TCCAGATGAT	AGTTTGACAT	TGCACACGGA	CTTGTACCAAG	ATCAACATGA	TGCAGGTTA	1320
CTTTGACCAA	GGGATTACA	ATAAGAACGC	GGTCTTGAG	GTGTATTCTC	GCCACAGGCC	1380
TTTTAAGAAC	GGCTATGCGG	TTTTGCAGG	TTTAGAAAGA	ATTGTGAACT	ATCTTGAAGA	1440
CTTGCCTTT	TCAGATAGTG	ATATAGCTA	TTTGGAGTCG	CTTGGTTATC	ATGGGGCGTT	1500
CTTGGATTAC	CTTCGCAATT	TCAAGTTGGA	GTTGACCGTT	CGTTCTGCC	AAGAAGGGGA	1560
TTTGGTTTT	GCTAATGAAC	CGATTGTGCA	GGTGGAAAGGA	CCTCTAGCCC	AATGTCAGTT	1620
GGTCGAAACG	GCTCTTTGA	ACATCGCAA	CTACCAAGACT	TTGGTGGCGA	CGAAGGCAGC	1680
TCGTATTCGT	TCGGTTATCG	AAGATGAACC	CTTGATGGAG	TTTGGGACAC	GTCGGGCTCA	1740
AGAAATGGAT	GCGGCCATCT	GGGAACACCG	CGCAGCTGTG	ATTGGTGGCG	CCAATGGAAC	1800
CAGCAACGTG	CGTGCAGGTA	AGCTCTTGA	CATTCTGTT	TTGGGAACCC	ATGCCCATGC	1860

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CTTGGTACAG	GTTTATGGCA	ATGACTATGA	AGCTTCAAG	GCTTACGCTG	CGACCCACAA	1920
AAATTGTGTC	TTTCTTGTGG	ATACCTATGA	CACCCTTCGC	ATCGGTGTAC	CAGCTGCCAT	1980
TCAGGTGGCG	CGTGAGCTGG	GTGATCAGAT	TAACTTTATG	GGTGTGCGGA	TTGACTCTGG	2040
GGATATTGCC	TACATTTCTA	AGAAAGTCCG	TCAGCAACTG	GATGAGGCTG	GATTTACAGA	2100
GGCTAAGATT	TATGCTTCTA	ATGATCTAGA	TGAAAATACC	ATCCTTAACC	TCAAGATGCA	2160
AAAGGCCAAG	ATTGATGTCT	GGGGTGTGGG	TACCAAGCTG	ATTACAGCCT	ATGACCAGCC	2220
GGCTCTTGGG	GCGGTTTACA	AGATTGTTGC	AATCGAAGAT	GAAACTGGTC	AGATGCGCAA	2280
TACGATTAAG	CTGTCTAATA	ATGCTGAAA	AGTTTCTACG	CCAGGTAAGA	AGCAGGTGTG	2340
GCGCATTACC	AGTCGTGAAA	AAGGCAAGTC	AGAAGGCGAC	TATATCACCT	ATGATGGTGT	2400
GGATATTAGC	GACATGACAG	AAATCAAGAT	GTTCATCCG	ACCTATACAT	ACATCAAGAA	2460
GACGGTTCGT	AATTTGATG	CCGTTCTCT	CTTGGTGGAT	ATCTTCAAAG	AAGGAATATT	2520
AGTTTACAAC	TTGCCTAGTT	TGACTGACAT	TCAGGATTAT	GCCCCTAAGG	AATTTGACAA	2580
GTTGTGGGAT	GAGTATAAGC	GTGTGCTCAA	TCCGCAGCAC	TATCCAGTGG	ATTTGGCGCG	2640
TGATGTATGG	CAAGATAAGA	TGGACTTGAT	TGATAAGATG	CGCAAGGAAG	CCCTTGGTGA	2700
AGGAGAAAGAA	GAATGAGTTT	GCAAGAACG	ATTATCCAAG	AGCTGGGTGT	CAAACCAGTG	2760
ATTGATGCC	AGGAAGAAAT	CCGTCGTTCT	ATTGATTTCT	AAAAAAGATA	TCTGAAAAAA	2820
CATCCCTTCC	AAAAAACCTT	TGTACTAGGG	ATTCTGGGG	GACAAGACTC	AACTTGGCA	2880
GGACGTTGG	CGCAATTAGC	TATGGAAGAA	CTGCGAGCTG	AAACGGGAGA	CGATAGCTAC	2940
AAATTATCG	CTGTCCGCCT	GCCATACGGA	GTGCAAGCTG	ATGAAGCAGA	TGCTAAAAAA	3000
GCCCTAGCCT	TCATCCAGCC	AGATGTCAGC	TTGGTTGTGA	ATATCAAGGA	ATCAGCTGAT	3060
GCCATGACAG	CTGCAGTTGA	AGCGACAGGT	AGTCCTGTTT	CAGACTTCAA	CAAGGGGAAT	3120
ATCAAGGCAC	GTTGCCGTAT	GATTGCTCAG	TATGCCCTTG	CTGGTCCCCA	TAGCGGAGCG	3180
GTCATTGGAA	CAGACCACGC	CGCGGAAAT	ATCACAGGTT	TCTTTACCAA	GTGGGTGAC	3240
GGCGGTGCAG	ATATTCTCCC	TCTTTACCGC	CTCAATAAAC	GCCAAGGAAA	ACAGCTTTG	3300
CAGAAACTTG	GCGCAGAGCC	AGCCCTTAT	AAAAAAATCC	CAACGGCAGA	CCTAGAAGAA	3360
GATAAACACAG	GCCTAGCTGA	CGAAGTCGCA	CTTGGAGTCA	CCTACGCAGA	GATTGACGAC	3420
TACCTAGAAG	GCaaaACAAT	CAGCCCAGAA	GCTCAAGCGA	CCATTGAAAA	CTGGTGGCAC	3480
AAAGGCCAAC	ACAAACGCCA	CTTACCCATC	ACCGTATTTG	ATGACTTTG	GGAGTAAAAA	3540
GGTCCGGGGG	ACCTTTTAG	CTTCTTGCCT	TGAAATTAAA	AAGCAAGAAA	AACTCCACT	3600

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GGAGGTTTTC AGCCTCTCAT CTTGAAATAA GAAAGTGAGA GAAGGTCTGG GGGATCTTGA	3660
ACCCCGAGTT TAGAAATAAG AAAATGAGGC AGATTCAAGTA ACTCGAAGAG TTCGATTCA	3720
TCGTCTTACC CCTGCAACGA TGACTAGGTT TGAAAAAGCT TGCTAGAGCG CATTCAAAC	3780
CAGGCAGCAA CTGCGTCAAG AAATTAGAAG ACAAACTCGT TTTCTAGCTG TTACTGAGTT	3840
GAGCCTTTT ACTACGAGTA TAGAAATAAG GAAGTGAGGT AGCATCATGA AATCTATCGG	3900
TACGCAAATA TTACAGACAG AACGTTTGAT TTTAAGAAGA TTTGTGGAGA GTGATGCAGA	3960
AGCCATGTTT CAAAATTGGG CTTCATCCGC TGAGAATCTG ACCTATGTTA CCTGGGATCC	4020
CCATCCTGAT GTCGAAATCA CTCGAAACTC GATTTGCAAT TGGGTTGCTT CCTATACTAA	4080
TCTCAACTAT TATAAATGGG CCATTGCT AAAAGAAAAC CCAGAGCAAG TAATAGGAGA	4140
TATCAGCATT GTTAAGATAG ACGAGGCTGA TTTAAGCTGT GAAATTGGCT ATGTGTTAGG	4200
CAAGGCTTAC TGGGAAATG GTATGATGAC AGAGACTTTG AAAGCTATCT TGGACTTTG	4260
TTTTACTCAA GCAGGTTTTC AAAAGGTCAG AGCACGTTAT GCCAGTCTCA ACCCAGCTTC	4320
AGGTCGTGTC ATGGAAAAGG CTGGAATGTC CTATCTACAA ACCATTGTTA ATGGTGTAGA	4380
GAGAAAAGGC TATCTTGCAG ATCTTATTAA TTATGGTATA AGTAGGGAAG AATGTTGAAT	4440
TCTATTTCT GTTTCTATCG AAGTCAACTA TTATGGTAA ATATAATAAT TAGCATTCCA	4500
AGTTTATTTG AAACTTAAA ATAGCATATT GATTAGTACA AGACAGATGT TCTAGTTCC	4560
TCTTTAATCT GGTTTAGTGT TAGTTAAAAA ATCGCTTTAA GCTTGTAAC AAGAGGGAGC	4620
TAATCGACTA GATTCTCCAG CCGAACAGGT GGTAAATGTAC TTTTATAGT GTAATCCTAG	4680
CTGTTGTTAA ATTTAAAATA GAATCCTCTA TCGAGTTAGG GAATTAAAATT CAACCAATT	4740
TATTCACTGTT TTTTCTATCA AATTATCTAA TATTAAAATA GTCTCATCT GATGAGAAAA	4800
CTATTCCCAA ATCATTCTATA CCTCTCTCAA CTAGATGTAA CTTACAAAAC CCCTGACCTC	4860
ATGAGCCACT TTCTTCCCTC TCATGAGGTC AGTTTTACTT TCTGCTGTC CAGTATCGTT	4920
TTTCCTCGCT AGATTTCCCTC AAAAGGGCAG ACTCCTCCCT TGGTGCCTCA CACGATTTTT	4980
TCATCTCGAC TGTTCTTAA TGCACTATTA ACAGACGTTT TCTTCTAGGT GGTCATAAG	5040
GAACAGGAAG ATTCAAGGTTG ACTTTCTAA TCCTAGAATA AAGTGCTGAA AACAAATTCGG	5100
AATAGGCATA GAGACTAGAC AATTTGAGGA GCTGCTTGC CG TCCTGTTCGA ACACATTT	5160
CCACCACGTG AAGAAAAGA TGGCGGAAGC GTTTGATTGT TAAAGTTGG AAGTCACCTC	5220
CAGCTAGATG TTTGAGAAAA AGATAGAGAT TGTAGGCGAT ACAGCTCATC ATCATAACGAA	5280
CTTCGTTTT GATTAAGGTT GAACT	5305

(2) INFORMATION FOR SEQ ID NO: 136:

919

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 3964 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 136:

TGGCAGCTCG TCGTCGAAA GGACGCAAAG	60
TTTGGCTGC ATAATCCAAA CGAATTCTAT	
CAAAAATCAG TAGGAACCTCG AGTCTACTGA	120
TTTTTATTTTG TGTAAGAAAG TTCAGTAGAT	
GCAAATGGAT TCGGAAGCGA TGTTACAGTA	180
GATTGAAACT AGAAATAGTAC ACCTCTGTT	
CTAAAACATT GTTAGAAATC GATTGACTG TCCTGATCGA	240
TTTGTCCTGT TATTATTTA	
TTTTACTATA AAGTTGAAGT AGGTTGGAGAT	300
GGTACAGCAA CAATCGTCTT TAAAGATGGT	
TCAGCTATTA CAATTCCAGG AAATCAATTG CTAGCACAAG	360
ATCCAAAAGC ACAAGATAGC	
ACTAAACTGA CTGCTGAAAA ATCAACTGTT AAAGCACCTG	420
CTCAAAGAGT AGATGTAAAA	
GATATAACTC ATTAAACAGA TGAAGAAAAA GTTAAGGTTG	480
CTATTTACA AGCAAATGGT	
TCAGCATTAG ACGGAGCGAC AATCAATGTA GCTGGAGATG	540
GTACAGCAAC AATCACATTC	
CCAGATGGTT CAGTAGTGAC GATTCTAGGA AAAGATACAG	600
TTCAACAAATC TGCGAAAGGT	
GAATCTGTAA CTCAAGAACG TACACCCAGAG TATAAGCTAG	660
AAAATACACC AGGTGGAGAT	
AAGGGAGGCA ATACTGGAAG CTCAGATGCT AATGCGAATG	720
AAGGCCGGTGG TAGCCAGGCG	
GGTGGATCAG CTCACACAGG TTCAACAAAC TCAGCTCAAT	780
CACAAGCTTC TAAGCAATT	
GCTACTGAAA AAGAACATCAGC TAAAAATGCC ATTGAAAAAG	840
CAGCCAAGGA CAAGCAGGAT	
GAAATCAAAG GCGCACCGCT TTCTGATAAA GAAAAGCAG	900
AACTTTAGC AAGAGTGGAA	
GCAGAAAAAC AAGCAGCTCT CAAAGAGATT GAAAATGCGA	960
AAACTATGGA AGATGTGAAG	
GAAGCAGAAA CGATTGGAGT GCAAGCCATT GCCATGGTTA	1020
CAGTCCTAA GAGACCAGTG	
GCTCCTAATG CTGCTCCTAA GACAACAAAGT GCACCGCAAG	1080
CAAATGCAGG AACAAATGCAA	
GATGTTACCT ACCAGTCACC TGCTGGCAAA CAATTACCTA	1140
ACACAGGTTA AGCATCAAGT	
GCAGCACTTG CTAGTCTTGG TCTAGTGGTG GCAACAAGTG	1200
GTTCGGCTTT GCTAGGAAGA	
AAGACTAGAC GTAGAAAATA GAACAGCTAG AAAATTCTAT	1260
TCTCTACTTA AAGTTAGATT	
ATAAGGGGGA TTTTGAGAAG TCATCAATCC TAGTGATGGG	1320
TGAGAAAAGT GAGAACCCAA	
GATAATCACA TACTTAGCT GAATAGGAAT ATTCTATCAA	1380
TGTAGCCAAT CTCTCTGTC	
TCTAACTGTG GAATAGGAGA TGGGCAATAT CGGATAGAAA	1440
AGATAGCAGA ATAGCTCT	

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ATTGAAGAGA GGAGGGAAA CCGAAAAATT AGGTGCCCT CCTCTTTTTT GGTATAATAG	1500
AAGATAGAAA ACGAGGTTAG AAGAGATGAT TTTTGATACA CATAACACT TGAATGTAGA	1560
AGAATTTGCA GGTCGTGAGG CAGAAGAAAT TGCCCTGGCT GCTGAGATGG GTGTGACACA	1620
GATGAATATT GTGGGTTTG ATAAACCGAC GATTGAGCAT GCCTTGGAGT TGGTAGATGA	1680
GTATGAGCAG CTCTATGCGA CTATTGGTTG GCATCCTACA GAAGCTGGTA CTTATACAGA	1740
CCAAGTTGAG GCTTACTTGT TGGATAAGTT AAAACATTCC AAGGTTGTGG CTTTAGGTGA	1800
AATTGGCTTA GATTACCATT GGATGACAGC GCCCAAAGAG GTGCAGGAGC AGGTTTTTCG	1860
CCGTCAGATT CAGCTATCTA AGGACTTGGA TTTGCCTTT GTTGTCCATA CCCGTGATGC	1920
GCTGGAAGAT ACCTATGAGA TTATCAAGAG TGAGGGCGTT GGTCTCGTG GTGGTATCAT	1980
GCATTCAATT TCAGGGACGC TTGAGTGGGC AGAGAAGTTT GTGGATCTTG GTATGACCAT	2040
TTCCTTCTCA GGAGTGGTGA CTTTTAAGAA GGCAACTGAC CTCCAAGAAG CAGCTAAAGA	2100
GTTACCTTG GACAAGATGT TGGTGGAAAC AGATGCGCCT TACTTAGCAC CTGTACCAA	2160
CCGTGGTCGT GAAAATAAAA CAGCCTATAC TCGCTATGTG GTCGACTTTA TCGCTGACTT	2220
GCGTGGTATG ACGACAGAAG AGCTGGCGGT AGCAACGACT GCAAATGCAG AACGAATTTT	2280
TGGACTGGAC AGCAAGTAAT GAAAGAGAAA ATTCTCAAG TTATCGTGGT TGAAGGGCGT	2340
GATGATACGG TCAATCTCAA ACGTTATTTC GATGTGGAGA CCTATGAGAC TCGAGGTTCT	2400
GCCATCAATG CTCAGGATAT AGAGCGGATT CAGCGCCTGC ACCAACGTCA TGGAGTCATT	2460
GTCTTTACAG ACCCAGATTT TAATGGGGAA CGGATTCCGGC GCATGATCAT GATGGTCATT	2520
CCAACAGTTG ACCATGCCCT TCTCAAGCGA GATGAAGCTG TTCCCAAGTC CAAGACCAAG	2580
GGGCGTTCTC TGGGAATTGA GCATGCCAGC TATGAAGACC TGAAACCGC TCTAGCTCAA	2640
GTGACAGAAC AATTGAAACA TGAGAGTCAG TTTGACATTA GTCGTAGCGA TTTGATTGCG	2700
CTTGGTTTTC TAGCAGGGGC AGACAGCCGT AAGCGTAGAG AATATCTCGG AGAGACTCTC	2760
CGAACCGCT ATTCCAACGG CAAGCAACTC CTCAAACGCC TAGAGTTGTT TGGGGTTACT	2820
TGCGAGAAC TGGAAAGAAC TATGAAATCT TATGAGTAGG AAAGATGTAG CCGTTACAAT	2880
TTTTAAAGTT TCACAGTATT TTTCGAAGCA GGTAGAAGAG GAGGCGTCTG ATGTTAATTG	2940
GTCAAAAAAT TAAAGAGATT CGGATAGAAA AAGGAATTAG TCGTCCAGAT TTTGTGGAG	3000
ATGAGCAAGA ACTGACAGTT CGTCAACTGT CGCGAATTGA AAGTGGAGCT TCGCAACCGA	3060
GTTTGCCCAA GTTAGACTAT ATTGCTCGCC GGCTAGGAGT TCCAGTTTAT AGCCTTATGC	3120
CGGATTTTC AGCTCTCCCT TCTGCTTATT TAGAATTGAA ATACCAAGATT TTACGTGAAC	3180
CAATCTATGG TAAAGAAGAG GAGTACGATA AGAAGGAAGC GTGTTGGAA GAGATTATA	3240

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AAACATACTT TGATAATCTT CCTAAAGAAG AACAAATTAGC ATGTGAAGTA TTGCAGCGT	3300
GTTGGATAC TTCTAGAACT AGAAGGCCTG AATATGCAGA GTTAATACTT GAGGAACATA	3360
TGCCTCAGAT TATAGAAAAA GAAGCTTATT CAATAAATGA TATGTTGTTG ATTCCGTTGT	3420
TTTTTTATCA AATGCTCATT AGAAAAGATC TTGCCAAATT TATAAATCAA ATCGAAAAGC	3480
TAATGCTCTT TCTTTGGAA CAGAAGAAGG TAACTCAAAT AGAGAATTAC TTTATAATTA	3540
GAGATACTCT TATTCAGGA ATGTGTTGTC TTGAAAAGGT AGGAGTAACT GATTGTTTA	3600
ATGATTATCT ATCGTGTAA CAAGAAATTA TGGATAAAAC TCAAGATTAT CAAAAGAAC	3660
CTCTTGTATT TATGTTTTG TGGAAAGCAAG CATTAAGAGA AGAAAGAGAT TTTAGTTAG	3720
CTGAATCATT TTATCAGTCT TCTAAACAT TTGCGCAGCT AATTGGAGAT GAATTCTAG	3780
TAAAGAAAATT GACAGAGGAA TGGCAAGAGG ATGTCAAAAA ATATTTATAA ACATAGTGAA	3840
TCAGTGACAA AGATGTCCTT GTCCCTCGTAT CAAAACAGTT CTAAAGTTCG TCTTAGGGA	3900
TGTTTTTTA GATATAAGCT AAAAATGACA CGAAATGGTT AGATTTAAG GACATTGATG	3960
TCCG	3964

(2) INFORMATION FOR SEQ ID NO: 137:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 12666 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 137:

TGAGACCGTT ATTTGTATTA GGGAAATGGG TATCTATTTT TAATGCTGTG GGGATTTGA	60
TTGTTTCTAT TATTCAAACC AAAAGCTTGT CAGGTATTGG AGCAGGATTG TTTAATCTAT	120
ATAACATTTTC ATCTTATATA GGTGATTTAG TTAGTTCAC TCGATTGATG GCATTAGGAT	180
TATCTGGAGC AAGTATAGCA TCAGCTTCA ATTTAATTGT TGGTTGTTT CCGGAAATAT	240
TGGCTAAACT GACAATTGGA TTAGTATTAT TCATTCTTT ACATGCGATC AATATTTTC	300
TATCGTTACT ATCAGGATAT GTTCATGGAG CACGTCTGAT ATTTGTTGAA TTTTTGGTA	360
AGTTTTATGA GGGTGGAGGA AAACCATTTC AACCTTGAA GGCTTCTGAG AAATATATTA	420
AGGTTATTAC AAAGAATTAA TGGAGGATAT ATATAATGGA ACATTTAGCA ACTTATTTT	480
CAACCTATGG AGGAGCTTTC TTCGCTGCAT TGGGAATTGT ATTGGCGGTT GGATTAAGCG	540
GTATGGGTC TGCTTATGGA GTTGGTAAGG CTGGGCAATC TGCCGCAGCT TTACTGAAAG	600

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AACAGCCTGA	AAAGTTGCC	TCAGCTTGA	TATTGCAATT	ATTGCCCGGA	ACACAAGGAT	660
TATATGGTT	TGTTATTGGA	ATTTAATT	GGTGCAATT	AACTCCAGAA	CTTCCTTAG	720
AAAAAGCGT	TGCTTATTTC	TTTAGCTC	TTCCAATTGC	TATTGTAGGA	TACTTTCAG	780
CTAACGATCA	AGGAATGTA	GCAGTAGCGG	GAATGCAAAT	CTTGGCTAAA	AGACCAAAAG	840
AATTCAATGAA	GGGAGCAATT	TTAGCTGCCA	TGGTAGAAC	CTATGCAATT	CTTGCTTTG	900
TCGTATCATT	CATTTGACC	CTTCGTGTAT	AAGAAATAAA	TTTGCAATT	AAAGGAGGTG	960
TCTAAATGAG	CAATTTAGAA	AACTTACGAG	AGTCTGTTAT	TGAACAAGCT	CATGAAAAAG	1020
GGCGTATGAA	ATTATTGGAT	TCCAAAAGA	AGATTGATGA	TGAATTGAA	ATGCAAAAGT	1080
CGCTCATTAT	AAAGAAAAAA	GAAGCTGAAC	ATGAACGAA	GTTAAAAGAA	TTGCAACAGA	1140
AATATCAAAT	AATTTTCAAA	CAATTAAGAA	ATAAGGAACG	CCAATCAACG	TTAGTATCAA	1200
AACAGAAAAT	ATTAAAAGAA	CTTTTCAAT	CTGCTTACT	AGAAATGGAA	TCTGGAGTG	1260
CAGATAAAAGA	AATGGAGTTC	ATCTATCGAA	TTCTGGAACG	ATATTCACAA	CAAGAGGTCA	1320
TAGTAACCTT	TGGGGAACGG	ACTTAGCTA	AATTCAATT	GGAACAATT	GAGAAATTGA	1380
AATTCTCTTT	TCCAAATTAT	TTATTAGTG	AAACACCTAT	CTCAAATGAA	TCAGGCTTAC	1440
TTATTCAAT	AGGTAAAATT	GATGATAACT	ATTTGTATAA	AACATTAATT	GGATCGATTT	1500
CTAAGGAAGA	AAGTTCAAGT	ATCGCAAATC	AAATTTTAT	CAATTAAGGA	TGAAATTGGT	1560
TAATCCTTCT	TAGAAATTG	GAGTATTCCA	ATAAAATTAG	AAAGGTATT	TATGGATACT	1620
AATCTTTTT	CAAAAATAAA	TACGACGATT	TCGGTAAAG	AAAACGATT	TATTACAGAA	1680
GAAAATTTC	AAAAAATTAT	ACAATCCAA	GATACGGAGA	CATTGGCATT	TATCTTAGAA	1740
TCAACTCCCT	ATCATTTATC	GATTGACATC	TTAGAAGATC	CTAGTCAGAC	AGAGATTCG	1800
CTAATGACAA	AATTAGTCAA	TGATTATAGA	TGGCCTATG	CTGAAAGTCC	GTCTGATATA	1860
ATTGTGACTT	TATTTGCTTT	ACGATATGTT	TATCATAATA	TCAAAGTTT	ATTAATTC	1920
AAGGCGGCAA	TTAAGAAAGA	TTTTCTAAA	TTATTAATTC	CAATAGGGAT	TTTTGATATA	1980
GAAAAGTTAA	AACATTAGT	TTCTCCCTTA	CATTCAGATA	CACTTCCTGA	TTTTATGGTT	2040
CGTGAAGTAG	AATCAATTG	GAATGAGTAT	GAAACTTTA	ATAATATTG	TGTACTTGAT	2100
GTCGGAGCTG	ATCTAGCATA	TTTTAACAT	CTGAAACTTT	TATCTAATGA	GTAGATGAG	2160
GTACTGTCTC	AGGTATTGT	CGAAATGATT	GACTTTATA	ATATTATTAC	TGTAAAACGT	2220
GGTTTATCTC	AAAATAAGAG	TCATGGGGAT	ATTTTACAAT	TACTTCAGA	TGAAGGAAGT	2280
ATTTCTGCTA	AAGAATTAT	ATACATTGTA	AAAAATCAAG	AAATATTGT	GTGGTTCAAT	2340
AAAATAAAATC	CAAGCTTAGA	TTCAATCTTT	TCAACTTATG	AATTGAAGAT	GCAGGACGCA	2400

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ACAATTCAT CTTCTGAGTT AGAATTTTA TGTGATTTAC TATTGTATAA AACTTTAGAT	2460
CAAGGAAGGT ACAATGTAGA GGGGCCGTTA GTTCTTGCTA GATATTTATT GGGATGTGAG	2520
TTTGAAGTAA AGAATCTCG AATGATCATA TCAGCTCTC AAAATACAAT TCCCTTGAA	2580
TCAATAAAAG AAAGGATACG CCCACATTAT GGAAGCTAAT AAGTATAAAA TTGGCATAAT	2640
TGGTAGCCGT GATATTATT TACCATTAG CATGATTGGG TTTGATATAT TTCCTGCCTA	2700
CCAAGAACAA GAAGCTATAA ATACACTAAG AAAATTAGCT CAATCTGATT ATGGTGTCAAT	2760
TTATATCACT GAAGACATTG CTTCAATGAT ATTAGATACA ATTCGCCATT ATGATTCCCA	2820
AGTTGTGCCT GCTATTATTT TATTACCGAC TCATAAACAA GGTTAAATT TAGGATTAAA	2880
ACGTATAGAG GATAATGTAG AGAAAGCAGT AGGACACAAAT ATTTTATAAT AATGTACAAA	2940
ATTGTCTGTA ATATTATTCT ATAATTTTG GACTTAGTAA GGAGAATAAC TTTGACTCAA	3000
GGGAAGAGTTA TAAAAGTATC GGGACCTCTA GTTATTGCAT CAGGTATGCA GGAGGCTAAT	3060
ATTCAAGATA TTTGCCGTGT AGGTAAGCTA GGGTTAACG TGAAATTAT TGAAATGAGA	3120
AGAGATCAGG CATCTATCCA AGTCTATGAA GAAACATCTG GTCTTGGTCC GGGAGAACCT	3180
GTTGTTACAA CTGGAGAACCT TCTCTCGTT GAATTAGGGC CAGGATTGAT TTCTCAAATG	3240
TTTGATGGCA TACAACGCCCTT ATTAGATCGA TTTAAATTGG CTACTCATAA TGATTTCTA	3300
GTTCGTGGGG TAGAAGTTCC AAGTTGGAT AGAGATATTA AGTGGCATTT TGATTCCACT	3360
ATAGCAATTG GTCAAAAAGT GAGTACGGGT GATATTCTT GAACTGTCAA GGAAACCGAG	3420
GTAGTTAACATC ATAAAATTAT GGTCCTTAT GGAGTATCTG GAGAAGTCGT TTCTATTGCA	3480
TCTGGCGATT TTACAATTGA TGAAGTTGTA TATGAAATAA AAAAATTGGA CGGTAGTTTC	3540
TATAAAGGAA CGCTTATGCA AAAATGGCT GTCCGCAAGG CGCGTCCTGT TTCTAACGT	3600
TTAATTCCAG AAGAACCAATT AATCACAGGT CAACGAGTTA TTGATGCATT CTTTCCAGTA	3660
ACCAAAGGGG GAGCTGCAGC AGTTCTGGA CCGTTGGAG CAGGAAAGAC AGTTGTACAA	3720
CACCAAGTAG CTAAATTGCA CAATGTTGAT ATTGTTATTT ATGTCGGTTG TGGAGAACGT	3780
GGAAATGAAA TGACGGATGT ACTGAATGAG TTTCTGAGT TGATTGACCC TAATACCGGA	3840
CAATCAATTG TGCAACGGAC AGTTCTGATT GCTAATACCT CAAATATGCC TGTTGCTGCT	3900
CGTGAGGCTT CAATTTATAC AGGAATTACG ATGGCTGAGT ATTTTCGTGA TATGGGCTAC	3960
TCTGTCGCCA TTATGGCTGA TTCAACTTCA CGTTGGGCAG AAGCGCTACG TGAAATGTCA	4020
GGACGTCTAG AAGAAATGCC TGGTGATGAG GGTTATCCTG CTTATCTGGG AAGTCGTATC	4080
GCTGAATATT ATGAAAGAGC AGGACGTTCT CAGGTTCTAG GGCTTCCAGA ACGTGAAGGA	4140

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ACGATTACTG CTATTGGAGC TGTATCGCCA CCTGGTGGAG ATATTCAGA ACCAGTTACT	4200
CAAAACACTT TACGGATTGT GAAAGTTTT TGCCCCCTTG ATGCTCCGTT GGCACAGCGA	4260
CGTCATTTTC CTGCAATTAA CTGGCTTACA TCTTATTACAC TATATAAAGA CAGTGTGGC	4320
ACTTTATATAG ATGGTAAAGA GAAGACAGAT TCCAATAGTA AAATAACTCG TGCGATGAAC	4380
TACTTACAAC GGGAACTTAG TTTAGAGGAA ATTGTTCGTC TTGTTGGAAT TGATTCTCTG	4440
TCTGATAATG AACGACTAAC GATGGAAATT GCTAAACAAA TTCGAGAAGA TTATTTGCAA	4500
CAGAACGCTT TTGATTCCGGT AGATACATTC ACTTCGTTG CAAAACAAGA AGCAATGCTA	4560
AGTAATATTC TCACCTTTGC TGATCAGGCA AATCATGCTT TAGAGTTGGG TTCTTACTTT	4620
ACAGAGATTA TGGAAGGTAC CGTGGCAGTT CGAGACCGTA TGGCGAGAAG TAAATATGTT	4680
TCAGAAGATA GATTAGATGA AATCAAAATT ATATCAAATG AGATTACACA TCAAATTCAT	4740
TTGATATTAG AAACAGGAGG TCTATAAATG AGTGTATCAA AAGAATACAG AACTGCTAGT	4800
GAAGTTGTTG GGCCTCTTAT GATTGTTGAA CAAGTAAATA ATGTGTCTTA CAATGAGTTA	4860
GTTGAAATTC AACTTCATAA TGGAGAAATT CGTCGTGGAC AAGTTTTAGA GATCCACGAA	4920
GATAAAGCAA TGGTTCAGCT TTTTGAAGGA TCTAGTGGAA TAAATTTAGA AAAGTCTAAA	4980
ATTCGTTTG CTGGTCATGC ATTAGAATTG GCTGTATCTG AGGATATGGT TGGTCGTATT	5040
TTTAATGGGA TGGGAAAACC AATTGATGGT GGACCAGATT TAATTCCAGA GAAATATTTA	5100
GATATTGATG GTCAAGCTAT TAATCCTGTA TCTAGAGATT ATCCAGATGA ATTATTCAG	5160
ACAGGGATCT CCTCTATTGA TCATTTGAAT ACTCTTGTAC GTGGTCAAAA ATTACCAAGTA	5220
TTTCAGGTT CGGGCTTACC TCATAATGAA TTAGCTGCTC AGATAGCAAG ACAAGCGACT	5280
GTTTTAAATT CTGATGAAAA TTTTGCCTT GTATTTGCAG CAATGGGTAT TACTTTGAA	5340
GAAGCTGAGT TTTTTATGGA AGAACTCAGA AAAACAGGAG CGATCGATCG TTGGTTTTA	5400
TTTATGAAC TGGCAAATGA TCCTGCAATT GAGCGTATTG CAACTCCCCG CATTGCTTTA	5460
ACTGCAGGCAAG AGTATCTAGC TTTTGAAAAA GATATGCACG TTCTAGTTAT CATGACGGAT	5520
ATGACTAACT ATTGTGAAGC GTTACGTGAA GTCTCGGCAG CTCGCCGTGA AGTTCCAGGG	5580
AGACGAGGCT ATCCGGGATA TTTATATACA AATTATCAA CTCTATACGA AAGGGCTGGT	5640
CGCTTAGTTG GTAAAAAAGG TTGGTGACA CAGATTCTA TTTTAACAAT GCCAGAAGAT	5700
GACATAACAC ATCCAATTCC TGATTTAAGT GGATACATTA CTGAAGGGCA AATTATTTG	5760
TCGCATGAGT TGTATAATCA AGGTATCGT CCACCAATCA ATGTTTTACC TTCTCTCT	5820
CGATTAAAAG ATAAGGGATC TGGAGAAGGT AAAACTCGTG GAGATCATGC TCCAACATG	5880
AATCAACTGT TTGCAGCTA TGCCCAAGGG AAAAAGTTG AAGAGTTAGC AGTAGTATTA	5940

925

GGAGAATCGG CTTTATCTGA TGTAGATAAA TTGTATGTGA GGTTTACAAA GCGTTTGAA	6000
GAAGAGTACA TAAACCAAGG ATTTTATAAA AATCGAAATA TAGAAGATAC GTTGAATCTT	6060
GGGTGGGAAT TACTATCAAT TCTTCCTAGA ACAGAGTTAA AACGTATCAA AGATGATTG	6120
CTTGATAAAAT ACTTACCTT GGTAGAAGTT TAATCCGGAA ATGGAGTGAT TATCTATGGT	6180
ACGTTGAAT GTAAAACCAA CTCGTATGGA ATTGAATAAC TTAAAGGAAC GTTTGACAAC	6240
AGCTGAACGT GGACATAAGT TATTAAAGGA TAAAAGAGAT GAATTGATGA GGCGATTAT	6300
TTCTTGATT CGTGAGAATA ATCAACTTCG GAAAGAAGTG GAAAGTTATC TAATTGATAA	6360
TCTAAAATCC TTTGCAGTTG CTAAATCATT AAAGAATTCT CAAATGGTGG AGGAATTATT	6420
TTCAATTCCA TCGAAAGAAA TTGAATTATT TGTTGAGAAA GAAAATATCA TGAGTGTAAAC	6480
AGTTCCCTAGA ATGCCATATGA ATATTACTTC TCAAAATGAG AACAGTGAAT ACAGCTATTT	6540
ATCTTCTAAT AGTGAATGG ATGATGTATT TGCTACAAATG AATAGTTAA TTATATAAATT	6600
ACTAAGACTG GCAGAACGTT AAAAACGTG TCAGTTAATG GCTGATGAAA TAGAAAAAAC	6660
ACGTAGACGT GTAAATGGTT TAGAATACTC GATTATTCCA AACCTGTCGG AAACTATTCA	6720
TTATATAGAA TTGAAACTAG AGGAGGCAGA AAGAGCCAAT TTAGTTCGTA TTATGAAAGT	6780
GAAGTAGATC CTTTATTAG ATTATTAATT AGATGAACAA ATATCAGCTT GGATAAGGCT	6840
TTAACGCTTT CTAAGCTTTT TTATTTGACA GTATCAGGAT ATCTTTTCGA AAATTTGGT	6900
TTGTTAGATA ATGAAAATGT TTCTACTAAT CTAGATTTAG GATTAGTAAA TCGTAAATGT	6960
AATTATATAG AAAGTAAGCG CGTCATAACA AGGTATCTAT CATTCACTGGA GCTCCTCCTG	7020
TATACTATTA GTAAAGTAAA ACTATTGGAG GATATTTAA TGCCACAAACC TATTGTCCT	7080
GTAGAGATTC CACAATCTCG TCGTTTGAT TCTAAAAGA GAAATGATAT TCTGCTTAAA	7140
ATTCGTATTG GCAAGCTTGA AGTAAGTTTT TTCAATCTC TCAATCTCGA AATGGTAGAA	7200
CAGCTTTGG ATAAGGTGTT GCTCTATGAC AATTCACTCTA TCTAGCCTAG GGGAGGTCTA	7260
TCTCGTGTGT GGGAAAATG ATATGAGACA AGGAATCGAT TCACTGGCTT ATCTGGTTAA	7320
AACCCACTTT GAATTGGATC CTTTCTCCGG TCAAGTCTTT CTCTTTGTG GTGGACGTAA	7380
AGACCGCTTT AAAGTCCTTT ACTGGGATGG TCAAGGATTT TGGCTACTAT ATAAACGCTT	7440
TGAGAACGGC AGATTGATTT GGCTAAGTAC AGAAAAGGGAT GTCAAAGCTC TCACACCAGA	7500
ACAAGTAGAC TGGCTTATGA AGGGCTTTTC TATCACTCCA AAAATATAGT AGATTGAAAC	7560
TAGAATAGTA CACCTCTGCT TCTAAAACAT TGTTAGAAAT CGATTTACT GTCCTGATCG	7620
ATTTGTCCTG TTCTTATTTCA ATTTTACTAT AAATCCATCA GAAAGTCGTG ATTTCTATTG	7680

926	
AAATGAGGAC TTTCTTTTA TACTCATCTG CTTCAAAAA GCATTCTAGT CCATCTCCGA	7740
TTAACGATGG ACTTTATCAC CTCCTCTCC AGTCCTGT A AACATCTTG GAGTTGATT	7800
ATGACATCTT CCAAAGTTA AAAGGTTA TTCTTAAATC CACGTTACG AATCTCTTC	7860
CACACTTGT CAATGGGTT CATCTCTGGT GTGTATGGAG GAATAAATGC AAAGCCAATA	7920
TTAGTCGGAA TCTTTAAGGT ACTTGATT TA TGCCATATAG CATTGTCCAT AACGAGTAAA	7980
AGATAATCAT CTGGATAAGC TTGTGAAATC TCCTATTCC AAAGCCCC TT TAGCGCATAA	8040
CTTTGGCTCA GCTTCTATTA TCGCTCACAC CATCCATCAG AAGTTAATC TGAAGGTACC	8100
CAATTATCGC CAAGAAGAAG ATTGGGCTAG GATGGGTTA CCAATCACAC GTAAGGAAAT	8160
CTCTAATTGG CATATCAAGG CGAGTCAATA CTATTTGGAG CCCCTTTATA ACCTCTTGCG	8220
AGAGAGACTA TTGACTCAGC CCTTACTTCA TGCGGATGAA ACTTCTTATA GGGTGTAGA	8280
GAGTGATAGT CAGCTGACTT ACTATTGGAC TTTTTGTCA GGTAAAGCAG AGAAACAAGG	8340
GATTACGCTT TACCACCATG ATCAGTGTG AAGTGGTCA GTAGTACAAG AATTCTAGG	8400
AGATTATTCT GGCTATGTGC ATTGTGATAT TTTGCGGCAG TAACTTAGGA CTTTAGTCCT	8460
CTAGTTCTGC CTATGCGATA GCAGTCCAAG GTTTAGGAGC AAGGCGACGC TAAGCTTGGT	8520
AAACTTCGAA CCGCTCGTCT GCTTATCGTC AACTGGAAGA AGCTGAACCTT GTTGGATGTT	8580
GGGCGCATGT GAGAAGGAAG TTTTTGAAG CGCCCCCC AGCAAGCGGA TAAATCATCC	8640
TTAGGAGCTA AAGGTTAGC TTATTGTGAT CAGTTATTTT CCTTGGAAAG AGACTGGGAG	8700
GCTTTGCCAG CTGATGAACG ACTACAGAA CGTCAAGAAC ATCTCCAGCC CTTAATGGAA	8760
GACTTCTTG CTTAGTGCCG GCGTCAGTCA GTTTAGCAG GTTCAAAACT AGGAAGGGCA	8820
ATTGAATACA GCCTCAAGTA TGAAGAAACC TTTAAGACCA TTTTGAAAGA CGGACATCTG	8880
GTCCTTTCCA ATAATCTAGC TGAACGCGCC ATTAATCAT TGGTTATGGG ACGGAGTAAA	8940
AGAGTCCAGT GGACTCTTT AGCCTAAGCT CAGTTAAAAA AAGCGAGGGT GGTTATTTTC	9000
TCAAAGTTT GAAGGAGCTA AAGCAAGAGC TATTATTATG AGTTTGTGG AAACAGCTAA	9060
ACGTCATCAA TTAAATAGCG AGAAATATCT ATCCTATCTT CTAGAATGTC TTCAAACGA	9120
GGAAACTCTC GTAAACAAAG AGGTTTTAGA GGCTTATTTA CCATGGACTA AAGTTGTACA	9180
AGAAAAGTGC AAATAAGAAA TCTCCAGATT AGGAACATATC CGTGAGTTCT CCAGTCTGGA	9240
GATTTTCAA TAGACTCCT GCGAAACAAA ATATGGTATA ATAGTTCTAT GAATGATGAA	9300
GCAAGTAAAC AACTAACCGA TGCACGATTT AAGCGTCTTG TTGGTGTCA ACGCACCGACT	9360
TTTGAAGAGA TGTTAGCTGT ATTAAAAACCA GCTTATCAAC TTAAACACGC AAAAGGTGGA	9420
CGAAAACCTA AATTAAGTCT AGAAGACCTT CTTATGGCCA CTCTTCAATA TGTGCGAGAA	9480

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TATCGAACTT ATGAACAAAT TGC GGCTGTT TTTGGTATT C ACGAAAGCAA CTTAACCGT	9540
CGGAGCCAAT GGGTTGAAGT AACTCTGTT CAAAGTGGTG TTACGATTTC AAGAACTCCT	9600
CTCAGTTCTG AGGACACGGT AATGATTGAT GCGACGGAAG TAAAATCAA TCGCCCTAAA	9660
AAAAGAATT A GCGAATTATT CTGGTAAAAA GAAATTCAC GCTATGAAGG CTCAAGCGAT	9720
TGTCAACAAGT CAAGGGAGAA TTGTTTCTTT GGATATCACT GTGAACATT GTCATGATAT	9780
GAAAGTTGTT C AAAATGAGTC GCAGAAATAT CAGACAAGCT GGTAAAATCT TGGCTGACAG	9840
TGGTTATCAA GGGCTCATGA AGATATATCC TCAAGCACAA ACTTCACGTA AATCCAGCAA	9900
ACTCAAACCG CTAACAATTG AAGATAAAAGT CTATAACCAT GCGCTATCTA AGGAGAGAAG	9960
CAAGGTTGAG AACATCTTG CCAAAGTAAA AACGTTAAA ATGATTCAA CAACCTATCG	10020
AAATCATCTA AACGCTTCGG ATTACGAATG AATTTGATTG CTGGTATTAT CAATCATGAA	10080
CTAGGATTCT ACTTTGCGAG GAAGTCTATT ATCAAAAATA CCATCAAGAT TATATAAGAT	10140
TGATACAGGA AAAGTTTAT TTGATGGTGT AAATATTAAT CAAATAGATA AAAAAATATT	10200
AAGTCAAAAT TTAGGAGTAG TTCCACAGGA TTCATTTTA TTGAACCGAA GTATTCTGA	10260
TAATATAACT TTAAAGCACG AAGTTACTTC ACAAAAGATA GAGGAAGTTT GTAAAGCAGT	10320
TCAAATCTAT GATGAAATCA TGGCTATGCC GATGAAATT AATACTATCA TCTCAGAGAT	10380
GGGGTCAAAT ATTTCAAGGTG GGCAAAGGCA ACGGATAGCA CTGGCACGTG CATTAATAAA	10440
TAATCCTAGT ATTGTAATT TAGATGAAGC AACTAGTGCA TTAGACACTA TTAATGAGGA	10500
AAGAATAACA AAGTATATAC AAAGTCAGGG CTGTACTCAA ATAATTGTAG CTCATAGATT	10560
GTCAACGATT AAGGATGCGG ATGTTATTT TGTAATGAAA GGTGGTAAGA TTGTTGAGTC	10620
AGGAAATCAT AAGTACTTAA TGGATCTTGG TGGAGAGTAC TACAGCTTAT ATACAAAAG	10680
GAAATGAGGT GTAAAGAAAA TGAAGAAAAGA AAATGAATAT GTAATTTAA CAACAGCCTC	10740
ACTAGGGGTG ATGATTGGAA TAGTGTTCG AATTTTTTA GATTTCCAG TTGAATATGG	10800
TATTTCTTTA GGCTTGTTGA ATGGAATAGT ATTGGTTCG CTGATTGTTT ACAAAAACAA	10860
TAAGAATTAA GCATAATTAA TTGCTGAAA CTAAGGAGTA GAGATGGCTA TAGTTGAAAT	10920
TATAAATCTA ACAAAAAGCT TTAAAGATAT TGAAGTTATT CATAACACTT AAATAATAGA	10980
GCAACTACAG TAGTAGCTTA AAAACATGAT TAAATCGCTA TTCTTAGGAG TAGCGGTTTT	11040
TCTTTTGTT TAATACTCTT TGAAAATCTC TTCAAACAC GTCAGCTTGT CTTTACCGTA	11100
CTCAAGTACA GCCTGCGGCT CGCTTCCTAG TTTGCTCTTT GATTTTCATT GAGTATAAAA	11160
AGGGTCAAGT AAGTATAGTA AATTGAAATA AGATATGAAC AAATCGATTA GAAAAGTCAA	11220

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ATTAATTCT AGAAATATGT TAGAAATTGG TTTGAATTCC GCAATCAATT TGTCAGTTT	11280
TTATTCATT TCATTTTATT TAATTAGATT TTCCAATTTC TTAATTCAAG CTAAAAATCC	11340
CCAATCGTAG TGATTGAGGA TTGAGTAAAT AAATCTAAA CAATACCTTG TGCAATCATG	11400
GCATTTGCTA CATTTCAAA GGCAGCAATG TTAGCTCCTG CAAGGTAGTC TTTATCAAGA	11460
CCGTATGTTT CTGAAGTCGT TTTAGCTGTG TTGAAGATGT TTGTCATGAT GTCTTGAGA	11520
CGGCCATCAA CTTCTTCACG AGTCATGAG AGGCGAAGAC TGTTTGCGT CATTCAAGA	11580
GCTGAAACGG CTACACCACC AGCGTTGGCA GCTTTGCAG GTCCGTAGAA GATACCATT	11640
TCTTTGTAAA CTTTGATGGC ATCAAGGTG CTCGGCATGT TGGCACCTTC AGATACACAG	11700
ATAACGCCTT GAGCAACCAA ACGTTTAGCT GCTTCACCGT TGATTTCGTT TTGAGTGGCA	11760
CATGGAAGAG CAATGTCATA GTTCCAGCG TAAGTCCATA CAGTACCTTC GTGGTAGGTT	11820
GCAGTTGCTT TTTCAGCTGC ATACTCAGTC AAACGAGCAC GACGTTTTTC TTTAACATCA	11880
ACCAAAAGAT CGAAGTCGAT ACCATTTCA TCGATGACAT AACCATTTGA GTCAGAAACA	11940
GAAATAACAG TTGCAACCGAG TTCAGTTGCT TTTGAAGAG CATATTGAGC AACGTTACCA	12000
GAACCTGAAA TAACGACTTT CTTACCAAGCA AAGCTGTTAC CGTTAGCTTT GAGCATTCT	12060
TCAGTATAGT AAACCAAACG GTAACCAGTT GCTTCTGGAC GAATCAAGCT ACCACCAAAT	12120
CCAAGAGGTT TACCAAGTCAA GACACCAGCA TCAAATTGGT TAAGACGTTT GTATTGACCG	12180
TAAAGGTAAC CAATTCACG TCCACCAACA CCGATATCAC CAGCAGGTAC GTCAAGTGAT	12240
GGTCCGATGT GTTTTGCAA TTCAGTCATG AAGCTTTGGC AGAAGCGCAT CACTTCAGCA	12300
TCTGTTTAC CTTTAGGATC GAAGTCTGAT CCACCTTAC CTCCACCGAT AGGAAGTCCA	12360
GTCAAGACAT TTTTAAAGAT TTGTCAAAT CCGAGGAATT TCAAGATCCC TTGGTTTACA	12420
GTTGGGTGGA AACGAAGTCC ACCTTTGTAT GGTCCAACAG CTGAGTTGAA TTGAACACGG	12480
TAACCACGGT TTACTTGAAT TTTCCATCA CGGTCAACCC AAGGAACACG GAAAGAAACC	12540
ACCGCGCTCAG GCTCAGTAAT ACGTGCCAAG ATATTTCTT CGATATACTC AGGGTGTGTT	12600
TCAAATACAG GTTCTAAAGT GTTGAAAAAT TCTTCAACAG CTTGGAGGAA TTCAGCCTCG	12660
TGCCGG	12666

(2) INFORMATION FOR SEQ ID NO: 138:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3083 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 138:

AGCAACTGTT	GTGAACCAAT	TCCGATAAAAT	TCCAAGAATT	GGTTAACAGA	GCCATTTGA	60
CAAAAAAATCC	CGATAAAAAGC	ATAGGCTTTA	AGGAGCAAAT	TGATCCAGGT	AGGAAGGATA	120
ATCAGCATGA	GCCAGAGTTG	ACGGTGTTCG	AGACGGGTCA	AAAAGAGGGC	CGTCGGATAA	180
CTGATAAGCA	GTGCCACAAA	GGTCACAATG	CCTGCATAAA	GCACTGAGTT	GAAACTCATT	240
TTAAGATAGG	TCAAGTTTG	TGACGCAAAG	TAAGATTGT	AATTTCTAA	ACTGAACGG	300
CCTTCGATGT	TGAAAAGGA	TTGACCGAAA	ATCAAGACCA	AGGGTGCCAA	TACAAAGAGC	360
GCAATCCAAA	GCATGTAGGG	TACTACAAAG	AGTTTAGAGC	TTGTTTCTT	CATCTCTTC	420
CTCCTCGATT	GCATTGATCA	AACCTGCTTC	TTGCTCTTCG	ATTTCTACGT	ACTCCTCAAT	480
ACGAGCATCG	AACTCTTCTT	CGGTTTCATT	GAGACGCATG	ATGTGGATGT	CTTCTGGTTC	540
AAAGTCCAGA	CCGATTTCT	CACCCACGAT	AGCCTTACGG	GTGAGTGG	TCATCCATT	600
ATTTCCAAGT	TCGTCATAGG	CGATAATTTC	ATAATGAAC	CCACGGAAAA	GCTGGTATC	660
GACCTTAACT	TGGAGCTTGC	CTTCTTCAGG	AAGGGTAATG	CGCAAGTCCT	CTGGACGAAT	720
AACGACCTCA	ACAGGTTCAT	TTGGCTTCAT	CCCACCATCA	ACCGCTTCAA	AGCGTTGCC	780
GTAAATTGCG	ACCAAGTAGT	CCTCAATCAT	GGTACCTGGC	AAGATGTTG	ACTCCCCGAT	840
AAAGGTGGCA	ACAAAGTGGT	TGATGGCTC	ATCGTAGATG	TCCACAGGGG	TTCCAGACTG	900
GACAATCTCG	CCATCATTCA	TAACGAAAAT	CCAGTCACTC	ATGGCAAGAG	CTTCTTCCTG	960
ATCGTGAGTG	ACAAAGACAA	AGGTAATGCC	CAATCGTTGT	TGTAATTCAC	GCAATTGTA	1020
CTGCATGTCT	GTTCTCAATT	TCAAGTCCAG	CGCTGATAAA	GGCTCGTCCA	ACAAGACCAC	1080
ACGGGGTTGG	TTGATGATAG	CACGGCGAT	GGCCACACGC	TGACGTTGTC	CTCCAGAAAG	1140
TTTGCAGGATG	GAACGTTTTT	CATAACCTTC	CAACTGAACC	ATCTTGAGAA	CTTCCGCTAC	1200
ACGCTGCTCG	ATTTCTTCTT	TATCAATTTC	ACGCAAGCGA	AGTGGAAAGG	CAACATTTC	1260
AAACACATTC	ATATGTGGGA	ACAAGGCATA	GGATTGGAAG	ACGGTATGTA	CGTCGCGCTT	1320
GTTGGTTGGA	ATATCATTGA	TACGAACACC	GTCAGTCATG	ATATCTCCTG	TCGTCGCATC	1380
CAGTAAACCT	GCAATAATGT	TTAGGATAGT	TGATTTCCCC	GAACCAGATG	CACCTAGAAG	1440
GGTGTAGAAT	TTCCCTTCTT	CCAACTCAA	GTTGATGTC	TTGAGAACCT	TGGTGTGCT	1500
GTCTTCAAAA	ACTTTAGAGA	CGTTTTGAA	TTCGATAATT	GGCTTTTCA	ATTGGCATAA	1560
ATTCCTTCTT	TTTCATAGAT	TAACCGATCG	GGGCTCTGTC	AGGTCCCCAC	TACCTCTTGC	1620
AGGGAGTAAA	ACCACCTGCA	TACATCTTCG	CTACCGATAG	GCTTCACCC	AAGATCCGGA	1680

930	
CTTCTCTTTC AAGCGTAATA CCTGAGTGTT CCTTGACTTT TTGATAACC GATTGGATCA	1740
AGTCCTCGTA GTCTTGCC GTTCCATCTG CGACATTGAT CATAAATCCT GCATGCTTT	1800
CTGACACTTC TACGCCACCG ATACGATAGC CTTCAAGCC AGCTTCTGAA ATTAACTGAC	1860
CTGCAAATG CCCGACTGGA CGCTTAAAGA CCGAGCCACA AGATGGGTAT TCCAAAGGTT	1920
GCTTGAGTTC ACGTAGGTGC GTCAAGCGGT CCATTTCTG CTTGATAACC TGATGGTTC	1980
CTGGAGCTAG GGCAAATTAA ACTGACAAGA CAACTGCACC AGACTCCTGA ATAGCTGAAT	2040
GACGGTAACC AAAAGCCAAG TCTTAGCGAC ACAGGGTTTC GATTTCTCCA TCCTTGGTCA	2100
AGACCTTACA AGACTGCAAG ATGTGAGCAA TCTGCCACC ATAGGCACCC GCATTCTAA	2160
AGACAGCACC GCCAACGCTT CCTGGAATAC CACAAGCAAA CTCAAAGCCA GTTAAACTAT	2220
GACGGAGGGC AATGCGAGTT GTTTCAATCA AGTTAGCCCC AGCTTCTGCT TCAATGGTAT	2280
AGCCATCAAC AGAAACGTTA TTGAGCTTGT CACACAAGAT GACAAATCCA CGAATCCCAC	2340
CATCACGAAC GATGATATTG CTTGCATTGC CAAGAACCAT CCAAGGGATA TTTTCTGGT	2400
TGGCAAATT CACAACGCGA GCCAACTCAA AACGATTCG TGAAAAGACC AAATAATCAG	2460
CCTCTCCACC TACTTTGTA TAACTATAGC TATGCAAGGG TTCCTTAAAA CGGATATCAA	2520
TTCCCTCTAA GATTCAAGC ATTTTTCTC TTACAGACAT GTCACTCTTC CTTTACAAA	2580
ATTCAATTCCA TTATACCAATT TTTAGAGACA TTTGACGACC ATAAAAAATAC CTTGTTGG	2640
TTTGCATAA GAAAAAGAGG TTCCCCCTT TTTATGATT TTTACAAAAG ATTCCTTGG	2700
TTCCATAGGC GACCAGAACG AGCTCCAGTG CTAGAACATC TTCAACCAAG ACTGGATTTG	2760
TCAACCAGCC TACTTGGAAA AGAGATGGTG CCAGATCAAA GAAGGCATGC AAGCCATAGG	2820
CTGCTAGGAG ATAAATCCAT TTCTCTGGC GAACAGCTTG GTAAACCCAA ACTGTCAAAA	2880
GTAATTGGAA ACCAAGCGCC AAGATCGCT CAAAACCAAG CAAATAAAATC TGCCAGACCG	2940
AAAGTGACTG AATGGTTTT AACATATTTT CAGACAGTAA TTGCAAAACC TGTGGATTCT	3000
GAGTTTGAAC TGCCGAAAGA ACAATGTAAA GATTGAGTAA ACTAGTAAGG CCTAGAAAAA	3060
TCAACTCCAA GCCACCAGTC CCC	3083

(2) INFORMATION FOR SEQ ID NO: 139:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 15363 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 139:

931

CCGGAGGATA TTGACCACCA CCAAAAGCAG GGGGAAAATC GAAATCAACC AATAGTAGGC	60
TACTGCGACA CTGGTCAACT CACTATCTGA TGCTTGATAA TAATGCAAAA AAGCTTTAA	120
TAAAGGTTTG TCTATCAGCT CTTTCCACCA CTTTTTCATG TCATACTCCT TCACTTATAA	180
TCTTATACTC AATGAAAATC AAAGAGCAAA CTAGAAAGCT AGCCGCAAGC TGCTCAAAAC	240
ACTGTTTGA GGTTGTAGAT AAGACTGACG AAGTCGATCA CATAACATACG GTAAGGCAC	300
GCTGACGTGG TTTGAAGAGA TTTTCAAGA GTATTAACTA ATTTCTTCTT ACCAATTCCA	360
CCATATCATA CGGTAGGGTA TTGGCAGCTT CCTTCAAGGA ATAGTTCTCT AAGTTATTAA	420
CATTTGTCG TAATTTCTTG GCATACTTAG TCGTAATCAA TCGTTTTCT TCGTATTCGA	480
AAATCAACTT GCGCTCCAGA TAATAGCCTC TCAGCATTTC ATCGATATTG TTGGGTTTGA	540
CACGATTGAT AACCCGTTCG ACAAAAGGCAC CACTGCTGAT AATAGCTGTT TCTCGAAGAC	600
GAGACTCCTG CATAAAACTA ATCAAAGAGC GTCTGTAGAC TCCCTTCAGG TTTTCCAAAC	660
TTTCAATAAT CATCTCTGTA TTGGCAAGAT AGAGCTCTGC AATTGGTCA TAATCAAGAG	720
CACGGAGAGC GCTTTGCTCC TTGTTCTCC AGCTACGGAA GGTCTTTCCG AGAGTAAAAA	780
CTTCATGAAG GAGAAAACGT AAAATCCTCA AGGAAACAAG AAAATAATAG GTCAGTCTG	840
AGGCAAGTTT ACGATTGATT CCTGTTCTA TATTTTCAG ATAACGTTGG TAAACTCGGT	900
AAGCACGATT GCTAATGTTCC CCTCTTCCAT AGGCCTGTT CAAACCATCA CTTTCAATAC	960
TAAGAACCAA GAGTTCAAA GCAGCCCAGT CTTCTTGATC ATCCTGGTTT TCTTGGCTTA	1020
AAATGAGATT TTCAATACGT CCATGATAAT TGTCAATAGC CGCATAGAGG GGAAGTTTAT	1080
TTCTGGTGTGTTTCCAACTCT TTTTCAACT CTAGCGTTAC TTCATTCAAA ATGGCGATAT	1140
GCATAAGATA ATCCTTGCTT TCTTCCCTTT CATCAGAAAG ATGAGGCAAG ACCAAGAGAC	1200
CTGTTAAAAA GCTAACAAAGC GTCACACCTG CAACAAGGAA AAGCAAAAGA GGATACTCCT	1260
GTTCTAGATT ACTTGGTATC AAGAGAATCG TAGCAATCGA CACCGTTCCC TTAACACCTG	1320
AAAAGGTCAA GAGAACATG TCCTTCATAT ACTTATTTAG CTTTTCTTG AGGCCTCGGG	1380
TTCTATAGGC ATAATAGCCA TAGATCATAA TAAAACGAAT GACAAAAAGG ACAAAAGGTA	1440
GGCGATAAG AGATAGCAAT AAAAGTAGAG GATTATAGAT TGGATTGGTC AAGATAGGTT	1500
CTGCTATCAT TTCCAACTCC ATCCCTAAAAA TCACAAAGAC AGAACCGTTG AGCATAAAGG	1560
TCACTGTATG CCAGACCGTC TCGGTACCG TATCCACTTG GGCTTCGAGG AGCGTGATTT	1620
TCTTGAAGCG ACTTGCCTTT AAAATTCCAG CAACTACGAC GGCAATAATA CCTGAAACAT	1680
GAACCTCTTC TGCCAGAAAG AAGGTCACTA GAGGCAAACCT CAATTCTAAT AAAAGTTCAC	1740

932

TGGCAATATC CGTTGCCGC ACACCTAGCA AGAAGGTATG GAGGAAGCGG TTGGTCATGG	1800
CTGTTAAAAA TCCAATTAAA AAACCGCCTA GGATTGAAAA GATGAGCGAA CTGCTAGCTT	1860
GCCCCAGAGA AAAAGCTCCA GTTGTCCAAG CTGTCAAAGC TACCTGAAAA GCCACCAAAC	1920
CAGAACATC ATTCAAGAGT CCTTCGCCCT TAAGAATATT GGACACGCCG TTAGGAAAGC	1980
TAAAACGCTC CGAAAGAGAG GCAAAGGCCA CCAAGTCCGT AGGACCAAGG GCTGCCCAA	2040
CAGCCAAGCA AGCTGCCAAG GGAAGGCTGA ACCAAAGAAG ATGGGCCAAG CCACCCAAAC	2100
TCAGGGTCGA GATAAAAATC ACTGGAAATA TGAGATAAAC AATGATTCGC CAGTGTGTTA	2160
AAATAGCCGT AACATCTGCT TCTTCAGCCT CTCGGAAAAG CAAGGGTCCG ATAACCAGTG	2220
CCAAAAACAA CTCCGTATTA AGGTGAAAGT CAGTATTGGG TAAAAAGAGA CCAATCACAA	2280
TTCCCAAAAG AATTTGCACC AAAGGGAGAG GCAAAAAGGG CAGGAGCTTA TTGGTTGTAC	2340
TTGAGACAAT CAAAACAGT AAAAATAGGA TGAGGTAAAT CAGTAATTCC ACGCACGTCC	2400
TCCTTAATCT TTTTTACAAC AGGATTCAAA TATCTCCTTC TGCTCTTGA TTTTTGGTC	2460
AATCTGGAA CAGTCTTGT GCTCAATTTC TCTCTGGCAC CGTCCATT CAAGAGCAAC	2520
TAATTTTTTC TTGATTTAA GCATTTTTT GCTCATATGC GCTTGGCTA GCACGCCAT	2580
CGCTCGTTCC TGGTGGGTTG ATTCAACAAA ATTCTGGCGC ATGGCATCCA GCTTTCGTG	2640
TAAGTATTGT TTATCCATGT CTGTATCTCT CTAATTTTC AATCATCACT AAAAACGGCG	2700
GGTTGTTGAC TTGGTTAAA GTTCCGTAAA TGGCAGCTGT GTACTCTTGT TGGTTCAACT	2760
GGATCACAAA ATCCAAGACA GCATCTCT CGAGATGCC TCCTCATGA CCATAGTAAA	2820
TCATAATAGC AATTCGTCCA CCTTGACAA GTAAGCCACA TAGCTTTCT AATGCCCTAA	2880
TCGTTGTCTG CGGTCGGGTG ATGACAGACT TATCAGCTGC CGGCAAATAG CCCAGATTAA	2940
AAATCCCTGC CTTAGCTTTT ATCACAAACT GGTCCAGTGT CTCATGGCCT TGCAAGATTA	3000
ACTGGCATT TGTCAAGTCA GCCTGATGCA AACGCTTTG GGTCTTTCC AAGGCTTGCT	3060
TCTGAATATC AAAGGCATAG ACTTGCTTGG CTAGCTTGGC TAAAAAAAGC GTGTCATGAC	3120
CATTTCCCAT AGTCGCATCC ACTACGACAT CCTCTTTGT CACGACCTCA GCCAAAAAAT	3180
CATGTGCCAT CTCAAGTGGT CTTTCATTT TCAAACCTCCT GTTTTACAGC CTTGCATCCT	3240
TGAACACTTC CACGACGTG CATCTCCATC TCAATGCTGT TGAGGACTTC CCATTTATTG	3300
AGGCTCCACA TAGGACCAAG CAGCATATCC CTAGGCGCAT CTCCGTAAAT TCGATGGATG	3360
ACGATATGTT TGGGAATAAT TTCCAGTTGG TCACAGATGA CCCTGACATA TTCGTCCTGA	3420
CTCATCAATT GTAAACGCC CTCATGGTAA TCTCGTTGCA TACGAGTATT TGTCAATAAGA	3480
TGGAGCAAAT GCAGTTAAT CCCTGAATA TCGTTATCCG TGACACAACG GCGGACATT	3540

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TCAACCATCA TCTCATGGGT TTCACCAGGC AAACCATTGA TCAAATGGGA AACAACTCTCA	3600
ATTTTTGGAT ACTTTCTCAA ACGCTTGACC GTTTCCACCT ACAATTCTATA AGAATGCGCA	3660
CGGTTAACCA GGTCAAGAGGT TGCTTCATAA GTAGTTTGCA AGCCCAATTC AACCGTCACA	3720
TGCATGCACT CCGATAACTC AGCCAAATAT TCGATGGTTT CGTCTGGTAA ACAGTCTGGG	3780
CGCGTTCCAA TATTGATTCC TACCACACCT GGCTCATTGA TAGCCTGTTCA ATAACGCTCT	3840
CGAATAACTT CCACCTTTTC ATGGGTGTTG GTAAAATTTT GAAAATAAAC CAGATACTTC	3900
CGAACATCCG GCCACTTGCG GTGCATAAAG TCAATTTCCT TATAAAATTG CTCACGGATA	3960
GGGCATCCG GTGCCACAAT GGCATCTCCA GAACCAGAAA CCGTACAAAA AGTACAGCCC	4020
CCATGAGCCA CAGTCCCATC ACGATTGGGA CAATCAAATC CCGCATCAAT AGGGACTTTA	4080
AAAGTCTTTT CTCCAAAGAG TTTTCGATAA TAATCATTCA AGGTATTATA AGATTCATG	4140
ACTTTCATTA TAACAAAAAT CACCCACAAT CTCAAAAGCC TGACTTTCTT ATAAATTCCT	4200
CTGTTTCTCG TTTCCATTAG CCTTTTTTTA TGATACAATA TGGGTATGAT TTTAATGAAA	4260
TTAGCATCTA TTTTATTATT GATACTGACC TTAGTCGTCT GCATTATCCT AACCAAACCTT	4320
TTTAGATTAA AAAAACTAGG ACGAAACTTT GCGGATTTGG CTTTCCAGT CTTGGTATT	4380
GAGTATTACT TGATTACAGC TAAAACCTTT ACCCATAATT TCCTCCCTAG ACTGGGGCTA	4440
GCCCTCTCGA TCCTAGCCAT TATTCTCGTC TTTTCTTCC TTTTGAAAAA ACGCAGCTTT	4500
TACTACCCTA AATTATCAA ATTCTCTGG CGTGCAGGAT TCTTATTAAAC CCTTATCATG	4560
TATATAGAAA TGATTGTTGA ATTGTTCTTA ATGAAATAGT CGAATCCCTA AGCATTTCCT	4620
AGGGATTTTT GCTTCTCTA CAAAATAGTA TAGACAATAA CACTATACAA TTTTATACAA	4680
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AACGTCACTT CTACACCTCC TATCATGCCT TTGATGCCCG GTCTTATCAC CCTAAAATGC	4980
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TGCCAACAAAG TATAGTTTG TGTGGAAGAA AACGACAGAG AAATTCTCCG CCAAACCTCA	5160
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GTGGATAAG AAACAAAAGA GAGGGTATAA AGAGTCAGCT AAAAACTTAG CGAATTGGCA	5280

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CGAACCTGAA TCAGCCCCCTC AAAAGGGACT GTATATGAAC GAACGCTATC AAAACTGAA	5460
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TCTGAGAGGG AAGCGTCAAG TGAGAATTGA CATGGGATTG GTACTTATGG CCAATAACCT	5640
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GAGAATGACT CAAAATTAAA AAGCTAGAGT TCCACAATTG GAAATATCTA GCTTTTTG	5820
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GGTCTTTAAT GATAAAGAACAG GTATCAAAT TTCTAGTCTT CTTTTTTACC TTTAGTAAC	5940
ACTAATCCTG CACTCAAACC TAGAAGAGTT AACCTGCTG CTACTGCTGC TTGGCTGCC	6000
GCACACCTG TACTTGGTAA CTGGCCTTA TTAGTTGAC TAGCTTCACT TGAATCAATT	6060
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GTTCCAACCA AGACGATGCG GTCTGTCGGA ACTTCTACCA CTTCACGGAG TTTTCTTCC	6180
TTACTTCCAT CAGGATTAAT CGCTGTAAAG ATACGTTCTT TTCCAACCTTT TCCTTCTTGT	6240
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TGAGTGCTTT CTTGAGGCAA TTCACTTGGT GCAAGGAAGG TCATCTCAAT CATCGCAACA	6660
CCGCTTTAT CTGCTTTACG CTCCACACGC CACTCTCATAG CTTGGCTTT GATAGCTTTA	6720
AATGTTACGT TGATTTCATC ACCAGCTGCA ATGTCTTAT CCGCACGATA AGAACAGCT	6780
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TCCGTACAC TATCTGCAAA GAACTGAACT TTTCTTGTG TAACAGTCCG TTCTACAATC	6960
TTACCATTCTT CACGGAAAAT CACACCCGCT GATACTTCTG GATTAGAAGA TGGTGTGGT	7020
GACCAGTTG TCCAACGACG ATTTTCTGAA TGATCTCCGT CATTGAGATA GTCAACGCGG	7080

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TCATGAGAGT	TTTGTCAT	ATCATTGGTT	GCTGAAGCAA	AGGCCTGGTT	ACTGTTTCA	7140
TCATAGTTAG	GGTTATCTGA	AAGAGTCTCA	CCAAGTTGT	CTGTCACTCG	TACAGTGATC	7200
TCAGCAACAA	GGTTACTACC	AAGGACACGG	CCTCGAACAG	TAAATTGACC	TGCTTTGTC	7260
AGATTTCCG	CTGGAACCTTC	TTCCCATTCA	ACTGTCAGGT	CTTTGTTTC	GTAGCCGTCT	7320
TTACCTGTGA	AGTAAACTGG	AACCTTAGTC	GGCAATTCAA	GTGCTTGACC	TACTTGTAGC	7380
AAGCGAGCTT	GTTTAACCGC	AGCAACTGGT	TTATGAGAAA	GTAAGCTCTT	ATCCTTAGTG	7440
AAGTGCAGAC	GGTATTCTCC	TAAGATGTCG	CCATTTTCAG	CTTCGCGAT	GACACGAAC	7500
GGCTCACCTT	CACGAACGCT	TGGAACGACG	GTAGCGAGAC	CATTGTTGCT	AACACTTGCT	7560
GTGACTGCCG	GAACTTTCC	ATCTACAGAC	TCAAGGTAGT	AGTCTGTCAA	ATCAGGGTTG	7620
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ACCATGCGAA	TACGAACAGC	ATAGGTTCA	ACTTTATCAA	AGCTAAAGTG	GTTCATTCT	7800
CCAGCCTTGA	GTTGAGCAGG	GGCTTTAGA	TTAGTAACTG	GTTCAGTT	GGCAGAACATCA	7860
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AATCCGACAC	TTAGATTATC	AACGGAGCGT	TTGCTCAAGA	TACCTGAATC	TCCAAACAGA	8040
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TCAGTTTGAG	CAGATACGCG	AACATGAAGT	TTAGTTGTTA	ATTGCGTAC	TTCTAACGGA	8280
CCATTAACG	TAAAGACACC	TTCCATTAGCG	TATTGCTCTG	GACGAATCG	ATCCCACATGCA	8340
ACCTTAGCTG	ATGAAACGTG	ACCATTGAA	TCATATGTCC	GAACACTTTC	TGGTAATTGT	8400
GGTGCTTCTG	CGATTGGAGT	TGTCACACTG	ACTTCTTCAA	CTGAAACGAT	ACCTTCTACA	8460
GAGACTTTG	CACGCGCTTC	AAGGTCAATT	CCTTCAACTT	TACCTAGTAC	TTCAAATGTT	8520
TGATAGGAGT	CTAGTTTTTC	TTTCGGAATA	GCTTGCCAAG	TGACTTTATG	AGTTTAGGG	8580
AAACCTTGT	CATACTCAAC	TGTTACTGTT	GCTGGAAGAC	TTGGTTCTG	ATGCAAATCT	8640
GTCACTACAT	TTACAGGACG	GATGGATTGC	GCAATCTTCT	TCTCAGTATT	GGCTTGGATA	8700
GTGAGTTCAA	CTTGGTCTTT	AGCTCCCTCA	TATTCAGCGT	TCAGAGTGAC	TGCTCCTGGC	8760
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GCAATTGGT GCGCTTCTTC AAGGAATTGA ATTGCATAGG TTTGAAGAGG GCCACCAC	9000
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ACAGCTGCAT TTTAGCACT TGCTGTGACT TCTGGCAACT TAGCTCCATA AGCAAGAGTG	9120
CGGTATTGCA TTGGTTTTG ACTAGTAAGA CCTGTTACTG CCTCACCA ACCGTTACA	9180
GTTGGTACTG CAGGTGCCGC AGGATTGCCT TCTTCTACCA CAAGGGTTGC ATGAATTGGT	9240
TGACCTTCTA AATAACCGGT CGCTGAAATA CGAGAACCTG GAATTGCTAA CTTAGCTTTA	9300
TCTTCTCGG CAATCTCCA CTTGTCACACT TCATACTCTT CAACACTTCC ATCAATCAA	9360
ACATAGGAAA CAGATTGTC TACAGAATTG AAGTCAGTAT TTGGAGCAAT ACGTTTCACA	9420
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GCCATACCTT TCACCGTTAC AATACCAGGC TTGCTCACAT CTACTGAAGA CCAGGGTTACA	9540
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AATGCTTTAC GAATCCAAGA ACCATCTGCT TGCGCCTTAT AGCGTTCACG GCTGGCTTGT	9840
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TCCAAGGTAC CTGGTTGATA GGCAACTTTC CATTCAAGAT AAAGTTCAATT AGCATTGCA	10140
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TTACCTTCTG AGTCAGCTAC TTTGGATGCT AATTCTTGT TTTCCAGTT CCAGTGAGGA	10320
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TGATTTGGT TGTGCCATGG TGTAGGTTCA CCAATATAGT CCGTACCTGT CCAGATAAAC	10500
TGTCCAGCAT AGCCAGCGTT GTCACGGTCA AAAGTCCATG AAGCGGTTGC TGTTTCCCC	10560
CAACCCACAC GATCATTCC ATAATCTGAC TGTTCTAAAT TACGCTCAGG TCCATTGCTA	10620

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TGTTTCAATT CACGTTCAAGG GCGATAGTAA CTTCCACGTG TACGGGTAGC TGAAGATGTT	10680
TCTGATCCAT AAATCAACCA TTTTGGATGC TTAGCTCTAA GGGCTTGTA ATTATCTTCA	10740
GAATAGTTAA ATCCAACAGC ATCGAGTTCA TCAGCAATT TCTCATGCC TCCGCTACCA	10800
TTACCGAAAC GGAATTATTC TGCTCCCATG GTAACATAGC GAGTCTTATC AACATCCTTG	10860
ATAAACTTAA CCAAACGTTT AACAGTTGCT AAAGAGTGGG CATCACCATT AGCTTCACCT	10920
ATTCATTAC CAATTGACCA CATGAAGATA GCAGGGTTGT TTTTGCCTCT TTCGACCATG	10980
GTACGTAGGT CAAAATCAGA CCATTTTCA CCTTTTCGAG CTTCTGGGTG AGTGGCATCT	11040
TTTCAAAGA AACGTCCATA GTCATAAGGT TTCTTGCCAC CATAACCACGT ATCAAAGGCC	11100
TCTTCCTGAA CGAGTAAACC TAGTTCTGCT GCGATTTGCA AGGTTTGCTC ACTAGCAGGG	11160
TTGTGGGTTG TACGGATGGA GTTAACCTCCC ATCTCCTTCA TTTGTTTGAG ACGGCGATAT	11220
TCTGCTTTAT AGTTTCTTC TGCTCCAAGC GCCCATGGT CGTGGTGCAG GGAACTCCA	11280
TGGAATTAA TACGTTCACCA ATTCAAAGAG AAACCTTCAT TTGGAGTCCA GTGATAGTAA	11340
CGGTAACCAA ACAAACTCTT CTTAGCATCA ACCAATTGAC CGTCACGGTA AACACGCGTA	11400
ATCAATTCTG ACAAGGCAGG TTTGTCATTT AAAACAGTCC AGAGTTTGG TCTTCAACT	11460
TCTAAATCG CATCTAGGCT TGTTGATTCA TGTGCTTTA AGGTACGACT CGCTGTACGA	11520
ACTAAGCCTG TTACAGCATG ACCACCTCGT TCAACGATT TGTGCTTTA AGGTACGACT CGCTGTACGA	11580
TGGTCTTTGT CGTCCGTATT GACGATTTG CTGGTCACAT GAGTTCAAC CTTGCCATGT	11640
TGTTGTTCTT CAAGTTTGG TGTTAAAATA GTTGTCCCAT TTTTCTCAAC ATGCACCTTA	11700
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TTGACATAAA CTTGAGAATC CATGTAGACG CCATCAAAG TAAGGCGAAC ATTTTCTTG	11940
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CCACCTTCAT TTTGTGCAGG AGATTCTGAG TCGAAATCGT TAAAGATACT CCAGTCATAC	12060
GGTAAATCTA ATTTTTCCA CGTAGATACG TCTGCATCAG GTTAAATGGC TTCTTAGAA	12120
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TGATTCACTT CTTCATTGT TACAGCTTTA GCATCTCCT TGAGCGGTT TTCTGATTT	12240
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TCACCTTGTT	CTGTCCTTTC	AACTATATTT	TTAGTTCCA	AAGCTTTATC	AGCCTTTCT	12420
TCTACTATCA	TTTTTCCTC	TTTAGGTTTC	TCAGCAGTAT	GAGTAATAAG	TGTTTCATCC	12480
GCATAAACTA	CAGATTCTCC	AGCTATATTT	CCTCCTAATA	AAACTGCACA	AGTCCCAATC	12540
ATTACTGAGC	AAGCTCCAC	AGCAAACCTA	CGAATGCTAT	AAACTCTTT	CCGATTCCAA	12600
TGGCCTTCC	CCATAAAAC	CTCCTTATAT	TATATTTAGT	GCAGTTAGCT	ACTACCAAAG	12660
CCCAAGTGGT	ATACATGGTA	TGACAACCTA	GTTCACAACAA	TTTACACTCT	GCGAAAATCC	12720
AATTCAAAC	TCGTCAGTGT	CGCCTGCCG	TAGATATGAT	TACTGACTTC	GTCAGTTCA	12780
TCTACAAAC	CAAACCATG	TTTGAGCTG	ACTTCGTCAG	TTTCATCTAC	AACCTCAAAA	12840
CCATGTTTG	AGCTGACTTC	GTCAGTTCA	TCTACAACCT	CAAACCATG	TTTGAGCTG	12900
ACTTCGTCAG	TCTTATCTAC	AACCTCAAAA	CTGTGTTTG	AGCAACCTGC	GGCTAGCTTC	12960
CTAGTTGCT	CTTGATTT	CATTGAGTTT	ATATTTATA	GGAGGCGATT	ATTTGCTTT	13020
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ACCAATTGTC	CAACCAAGAA	GTTCGAT	TGGCCTTCA	AGAGTAGAGT	GAGTAATCAA	13200
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GATACCTGCA	AGTGGCAAGA	TACCATTCC	AACTTTGAA	AGAACGACTG	CTTCAATCAA	13440
CATGATTGGT	GCAAGTACGT	TGGCACAAAGC	CCAGATTCA	GCACGACCAAG	CGATGAATGG	13500
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ACCTTGAT	AGTGGTTCTA	CGGCTGCGAT	GAACCATGAA	CCGATAAGTG	AGAAGAGTT	13620
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AGCATCAAAG	TCATATTAT	CAAGGCCTGG	GAAGAATT	TCAAAATCT	TATCCAAAC	13860
CATGATAACT	GGGTTCATCA	TGTAGTCAT	GTGAGTTGAT	GTCATTGGTG	ATGAACCTGG	13920
GGCGTTAAGA	AGGTCAACAA	ATGTAGGTTT	CATCAAGTCA	GAGTTGATAA	TTTCAACAC	13980
ACCGACAAGG	ACGATAGCTG	CTGTAGCAAT	AAAGAGTGAA	ACCCCTTGAC	TCACACCATT	14040
GTTATCAGCA	TACCATTAA	TCAAGAGACC	TGTGATAGAC	AAGTGCCAGA	TATCAAAGAT	14100
ATCGACATCA	AGTGTATCTG	TTTCCTTCAT	AGCTAGCATC	ACTATGTTGA	CAATCAAACAT	14160

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GATGAGCAAG AAGTATAGTG TCCAAGCAGA ACCCCAAGTG ATTGTAGCAA GTGGTGCCCA	14220
ACCAAACGTCG GTAATACTCA ATTGGATACC AGTGTGTTCA ACGAATTGTTG CTAGTGATGC	14280
TGAGAAAGCA GTGTTTAGCA TACCGATGAT AGCACCGATA CCTGTAAGAG CGATGGCAAG	14340
TTTGATACCA CCTTCAAGCG CTTTGGAGAA TTTCACTCCA AAAAGTAAAG CCAATACTGT	14400
CAAAATGATT AACATGATGA CAGGTCCACC CATTCTAAG ATGGGATTGA AAACCTTCC	14460
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AAGATGACCA TTTCCCTGTGA AGAAGTCAT TAACTGAGCA AGAATGTTG TTTGACTTGA	15000
ACTTGAATTA TTAATGATAA AGAAGAGTAG GGATACTTCT ACTTCCTTAT CAGGAGCTAT	15060
CATATTGTGA AAAGTTATTG GTTTTCTAA TCGAACAAACC ACCACTTTCT CAGCTAGATT	15120
ATGAACAATA TCTGTGTGAG GAATCGCTAC ATTTGGCAAG TCCTTTCCTA GAAATTCCAT	15180
ATCTAACCCA GTTGGAAATG ACTTTTCACG CGTGATCAAG GCTTCACGAT AAGTTGGAGT	15240
GACAATTCT CGTTCTTCCA ATAAAGTTGC AACCTGATCA AAGAGTTGTT CTTGACTATC	15300
CGCTTCTAAC CAAAACACAA GGTTTTGTC AAAGAAATAA TCTAATACCA TAAGTTTTC	15360
CGG	15363

(2) INFORMATION FOR SEQ ID NO: 140:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28882 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 140:

TAAGACTATT TAATAGTGGA GTGAAATAGG ATACGAACAA ATTGATTAGG AAAATCAAAT	60
GAATTTATAG AAATCTTTA GCAGTTATGT TATCCTATTC TAGTTCAAA ACGCTATAGA	120

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AGCAGCATTG TGCTAGTC k A GATTCA G T T TT ACTATACTAA AACGAGTAGC TTGAAATCAA	180
AAAACCCACC CTCACAGGCA GGTTTTATCT GTATTATTCA GCTAGATTAT GCTTTACCTT	240
CTGAACCGAA TACGTCGATA CGTTCTCAA CCGATGCTTG GATAGCTTT ACACCGTCAG	300
CCAAGAATT ACGTGGGT C G AAGAGTTTT TCTTGT C GTA TTCTGCTTCG TTTGCTTCGT	360
AGTCACGAGC AAATTTACGA GTTGC G T A G CGAATGCGAT TTGGCATTCT GTGTTAACGT	420
TAACTTGGC AACACCAAGT TTGATAGCTG CTTGGATT T G CTCATCAGGA ATACCTGATC	480
CACCGTGCAA TACGATTGGG AATCCTGAA GAGCTTCTGT CAATTTTGC AAGTGGTCAA	540
GGTCAAGACC TTCCCAGTTT ACTGGGTAAG GACCGTGGAT GTTACCGATA CCAGCTGCCA	600
AGAAGTCGAT ACCAGTTCA ACCATTGCTT TAGCGTCTTC GATTGGAGCC AATTCA C TT	660
TACCGATGAT TCCATCTTCT TCACCACCGA TAGTACCAAC TTCAGCTTCT ACTGAGATAC	720
CTTTAGCGTG TGCTTTTCA ACAACTTCTT TAGCCAATT A AAGGTTTCT TCAACTGGAA	780
GGTGTGAACC GTCAAACATG ATTGAAGTAT AACCAACTTC GATA C ACTCA AGTGCATCTT	840
CGTAGTGACC GTGGTCAAGG TGGATAGCTA CTGGTACAGT GATA C CCATT GATTCAACAA	900
GGTTAGCGAT CAAGTTGCGA GCAACTTTGT ACCACCCAT GTATTTAGCA GCACCCATTG	960
AAGTTGGAT CAAAAC T GG A GCTTTTTAG CTC T GCTGC GCGCAAGATA GCTTGAGTCC	1020
ACTCAAGGTT GTTTGTGTTA AATCCACCAA CTGCATAACC GTTGT C ACGG GCTGCTTGGA	1080
CAAATTTTC TGCTGAAACG ATTGCCATT T TATCAGGCCT CCTGTATATT TTTATGGTC	1140
ATCCCATT T A CATTGTCAT TTTATC A TT TTTGCCAAA AAATCTAGTT TTTCCCGCAG	1200
TTTCGATTGA TTTTCTCTA ACTCCATCTA TGAAACCC T TTCTCTCCCT AGTCTTGGAC	1260
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CTTGC G ACTG GCATTGTCAG AAAAAGACTG GGCTAGTTT TCTTGT C CCT CTGAGCTACA	1440
GCTATTT C CG ATAAAAAGTT CTC C CTCTCT TTCTCCAATT CGAACTAAGC CACCTGGAAC	1500
AGAGTGCTTA ATGGCATTG C TGATGAGATT AGAAAGAATC AACTTCATAA CTGATGGTT	1560
TAGATAAGCC TGCTGATGGG TCAA A CTATT GTCTATCTGG AGCTCTTTT CCTTGGCTAG	1620
CAAGGCATAA TCTTGACCA GATTTGCGT CATCTGGAGG AGGTCAATTG TTTCCCTATC	1680
ATCTCGCAAT TCCTGCACAG AAGAGAGGG A AAGTATCTGC AGAACATGGT GATTGAGTT C	1740
ATCCACAATC CCCAAGGCAA CTCCCAGATA CTGGTCTCTA TCCTTATAAC GACCGATATT	1800
CTCTCTCATA TTTTCGATTA GGATTTCAA ACTAGCCAGC GGTGTTTCA ATT C ATGAGA	1860
AGCTCCTCGT AGGAATT C GA CCTTCATCTT CTCCAGCTGG AGAATGGCTT CATTCTTT C	1920

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ATGCAAGTCC	GCAATAACAG	TCAAGAGATG	CTGGTAGAGG	CTATTGATTT	GTTCCCTTGAG	1980
ATTACCTATC	TCATCCTTAG	AATCCACGCG	CAATCGCACT	TGGGAATCCA	GGTCCATCAT	2040
CCGACGGGTG	ACCCGCTGTA	TTTCCAAAAT	CGGTGCAACA	ATAGTCCGAG	CGTAGATGTA	2100
GGCCACCAAA	AGGGAAATCA	GAAAGGGAGGC	CAGCAAGGTA	TAGGGAAGAA	ACTGGAGACT	2160
GATTTGCTCC	GCTTCCTTTT	GTAAATCCAT	GGAAGCTAGA	AACTGGAGAA	TCATAGTACC	2220
ACCGTCTTGC	GTTTTCACCT	CGCGCTCCTC	AATAAAGAGA	GAGGTTGTCT	GGCGGCTGT	2280
GTCCAGAGGA	AGACTGTCCT	TGACTTCTAA	CTTGTCTCG	GTCATCTCAC	CTTTGACGGT	2340
CCCCTTGATA	TCACTAGTCT	GGGAATACAA	GTCTAACACT	TGCTCGATAAC	TCTGCCATATC	2400
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ACTCAGATAA	GTCGAAGGAA	AAACAAAATA	AATAGCTAAA	TGAAGGCAGA	TAACCAGAAC	2520
ACTAAATATC	GAGAAGGTAT	AGATAAATAT	CTTTGCAAAT	AAACCTGTT	GTTTCATTTT	2580
CGCTCCAATT	TATAACCAAC	ATTGCGCACA	GTGAGGATAC	AATCCAAGTC	TAGCTTTTC	2640
CGCAATTCC	TGATATAAAC	ATCAATAACA	CGGTCAAAGG	GAACCTCATC	TGTCGCTTTC	2700
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CTGACCCCC	CACGGATCAT	CTCTTCATCT	TCTACAATTAA	AAATTTCAT	ACTTTAACTG	3240
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GCTGGCCTT	TGTCTGCAAG	CAACTGACCA	CTAGATAAAA	CGTTGTGAAA	TTCCTTCTC	3360
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TCCACTATTC	TTGAATAGAA	ACACAAGATG	CAATCTTAT	TCTAGACTCA	TTTTTTCAA	3540
TTTATTCAACC	ATCCAGCAAG	AGCTCTTTG	GTTGTTTCT	AAGGAGATTG	CTTGAAGCAA	3600
GCGCCATAAC	GAGAACCAACT	AGAACCAAGG	CAAGGACAAA	AATGATGATA	AAGTCTGATG	3660

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CACCAAGGTT AGAGGCTTGA GCCGCCTTAC TAGCCTGTTT GGCAACACCT GAAGTCACAT	3780
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AGCCCTTCTT ACGAATATCC TCTCCTTGAA AAAGGATAGA ACCTTCAACA GGACTATCTA	5460

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ATAATCCAAA CACGAAACCA GTCCACGTTT TTCAAGGACT GGTTTGATA TAGCACGTTT	7140
AAGTACCGAC TTCTGAGCTA CTATAGTAGA TTGAAACTAG AATAGTACAC CTCTACTTCT	7200

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CCCTAACCTTA AGCAATCCCC ATAATCGTCT CGATTTCTTC TTCCATTGCT TCCAGATAAT	7380
CACTCGTAGG CGAGTACGCA AGCGCTCATC TATGCTAGTG ACTATACTTT TCATATTAT	7440
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CCTTCTTCCC TGGCACGGTT GATAATCTTA TCATCCACAT CTGTAAAATT GGAAATATAG	10740

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GGCGGTGAGG GCGCTCAGAC TTTCACCTT TTTCATCATG CTCAGGTTT GGCGGACGAG	12660
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ACAGTGCCAA	AGATGAACCA	AACATAGCAG	CCATTGGTGC	AGATGCATT	TCATCATAAG	14400
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GTGGGAAGAAA	ATCCCCAGTT	GCCATT	TAGACATAAC	GGCAGCAGTC	AAGACAGTTG	14640
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CAATTAAC	ACGACCCGCA	AAAGTCGTTT	GAAACACTTG	TTTGCCATT	TTAACCCCCT	14760
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CATTCTGTA	GAAAATAGG	AAGGTGACGT	CGCACTCGAC	GAGTGTAGG	AAGCTTATCT	15000
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TGGATGGAGA AGTACACCTG CTACAGACAT AGCTAATGTA GATGTTACTT TTAATTGTTG	28200
TGATGCAGAA TAAGCTAATA ACAGCGGTAA GAAATAATAT GGAGCATCCC CAAAAAAATGT	28260
CAAAAAAGCA ATAGTCTGAG AATCTGATTG CAATATACCA AGCATTGGTA AAATGATTAC	28320
CAAGACTTTC AACATACCTC CCCCTAACAT TGCTGGAATG ATTGGAGTCA TGGAACCAGC	28380
GATATACTCA ATGATTCTTT CTAAAATATT CCCTTGTC CCTTGAACAA CTGAATCGGA	28440

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TTCAAAATTG CCAAGTTAA CGAATTCTT ATAATAATTA GCTACATCAT TACCAAGTAT	28500
AATTTGATAT TGTCCATTCT TTTTCATAAT ACCTATTACA CCTGGTATCT TCTTCACATC	28560
ATCATCATTG ACTAAATTAA CATCTTTAA TTCTAATCTT AAACGTGTTA CACAATGGGT	28620
AACTCTATTG ACATTTTTT CACCTCCAAT TACATCGAGG ATTTTTGTA CCGTATCTT	28680
ATAACTCATG GTATTCTCCT ATTCTATTAA TCTAAATTAA TTGTTAACG AGAATATGA	28740
GCCATCAAAT AAACTAATTC ACTAGAACGTC AGCAAATAAT TGTACTCCGT TTGTATAAAC	28800
ATTGCTACCT GTTCACCACA TTCATATTCT CTAGGATATT TATTTTCAT TAATGCTAAC	28860
AAGTCTTCAT CATCATCGTC GG	28882

(2) INFORMATION FOR SEQ ID NO: 141:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12835 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 141:

GCCTATGTCT TTTCAAAAA AATGCTTGAC TTGAGACGGG AACTAGGGAA GTCTAAAGGC	60
GGAAGGCATT GATTTATACT CTTCGAAAAT CTCTTCAAAC CACGTCAACG TCGCCTTGGA	120
TTATATATGT AACTGACTTC GTCGATGCTT ATCTACAACC TCAAAGCAGT CCTTTGAGCA	180
ACTTGCGGCT AGTTTCCTAG TTTGCTCTTT GATTTTCATT GAGTATTATA TTACTTTCTA	240
TTTGTAGGAG GTGGCTTATG AAGATTCCTC TCTTAACCTT TGCAAGGCAT AAATTTGTTT	300
ATGTCTTGCT TACTTTGCTT TTTCTTGCTT TGTTTATCG TGATGTTTG ATGACTTATT	360
TCTTTTTGA TATTGATGCG CCCGATCTAG CTAATTCGA TGGACAAGCA ATTAAAAATG	420
ACTTATTAAA ATCAGCATTA GATTTCGTA TTCTCCAGTT CAATCTAGGT TTTTATCAAT	480
CATTTATTAT TCCAATCATC ATTGTTTGCG TAGGTTTCA ATATATTGAG CTGAAAAATA	540
AAGTTTTACG ATTGAGTATT GGAAGAGAAAG TGAGTTATCA AGGGTTAAAA AGAAAGTTGA	600
CTTTGCAAGT TGCAAGTATC CCTTGTTGA TATATTAGT GACTGTGCTG ATAATTGCAA	660
TTATAACCTA TTTCTTGAGG ACTTTTCTC CTCTGGATG GAATTCTCTA TTTTCTGATG	720
GAAGTGGTTT ACAAAAGACTC CTAGATGGAG AGATAAAAAG CTATTGTT CTTACTTGTG	780
TCCTACTAAT CGGTATTTTC ATCAATGCAA TCTATTAAAC ACAAATAGTT GATTATGTGG	840
GGAATGTGAC TCGTTCGGCA ATCACCTATT TGATGTTCT TTGGCTTGGT TCTATGCTGC	900
TTTATAGTGC CTTGCCTTAC TATATGGTTC CTATGACGAG TTTGATGCCA GCTAGCTATG	960

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GGGATGTAAG TTTGATGAAA CTCTTTACTC CTTATATCCT TTATATTGTC CCTTACATGG	1020
TGCTTGAAAA ATATGAAGAT AATGTTAAG AATTTAACCA ATATTTGCT AAATAGAAAG	1080
ATTGTTTAC TACTCGTAT AGTCTGTATG ATGATTTGA TAAACCACCT ATTGTCAACA	1140
GCGGTTCAAA AGCAGGATGC TGTTATCTT TTCAAGAGAG AATTGATTTC AATTTTTCC	1200
TATAATGACT ATTCTGAAGC GAATTTAGAA ATCCCCAAC TATTGTTAAA CCTTCGCTT	1260
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CACTTGATTC GCTATCAATC AAGCTCCTTT TTCGATTATA CAAGGAAACG ATTGGTTGTC	1380
ATTCCTAAAT TTTTACTCA AGATTGTTT GTCTGGTTTC TTGGTTACT TCCTCTAGGA	1440
ATTCATTTCA AAACAGTCGC ACTTTCTTT TTACTTGCTC AGTTAATGAT GTTGTACTTA	1500
CTACTGTCTT ATCTGATAGC ACTGATTAGT CGGGGCGCTG GTTTTCCCTT TTTCTCTAT	1560
TTTTTAGCAT TTGTGGGACA AGAATGGATG ATGGATCATA TTGTAACAGT GTATTAGTA	1620
CTCTTAAGTT TATTAGTTAT GTTGATTGTT AGTCGCTTGG AAGAGAAATT TAAGAAAGGA	1680
TAAACGATGA GACTTGAAT TATAATGGA CAGAAAATT ATGGGAAAG ACCTATTTA	1740
AATCAGTTGA ATTTGGTGT TCAATCAGGA AAAATTATG GACTTAAAGG TGATAATGGA	1800
TCTGGCAAGA CGGTTCTTT AAAGATACTT GCTGGTTATA TTAAGCTTGA CAAAGGAAAA	1860
GTTCTTCAAG ATGGTAAAGT TTACGGGTA AAAATCATT ATATTCAAGA TGCAGGAATT	1920
TTAATTGAAA AAGTCGAGTT TTTATCTCAT TTATCCCTGA GAGAAAATT GGAACTGTTA	1980
AGGTATTTT CATCTAAAGT TACGGAAAAA AGAATTGCCT ATTGGATTCA ATACTATGAT	2040
TTACAGGAAT TTGAAGACAT TGAATACCGT CATTATCCT TAGGAACAAA GCAAAAATG	2100
GCCTTGATTC AAGCCTTAT TTCCCTCTCCT TCTATACTCT TTCTCGATGA ACCTATGAAT	2160
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GAAAATGGTC TGGTTATCCT GACGTCGCAC ATATCGGAAG ATATTCAGA CCTTTGTACA	2280
GATGTATTAG TTGTCGAAAA TGGACATATA CAAATGTAAA GGATATACAA TCCTAGGAGA	2340
TGGCTTATGG CACATCTAAA ATCATTTATT ACACGATATT CCAAGGTTA TATTGGTTA	2400
GTTCTGCTGA TCTGGCTGTC TTTCTTCTTT ATCCCTGGG ATAAACCACT TCTGGGGATA	2460
AGGATTGACA TCTTCATCAT ACAGAAAATC TTGCTAGCTT TTGGAATTCT GTCCATTCTC	2520
ATGGCCTTGC TGTCCAAGAA AGTCAGTCTC TTTGTTTTG GACTGATTG CTGTCTTCT	2580
CTTTGGATTA ACTTATTAT CACATTTGCC ATTTGCCGA TTTTGGCAA TTAAACAGTC	2640
ATAAAAGTCG GAGAGGTTAG CTTGAAAATC AACCTCTTT TCCTTTCAA AATGGGGATT	2700

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CTTCCTTGAA AATAATCAGT AATTGTGCTA AAATTAAAGG AACATTCTAA AATATTCGGA	2760
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TGGAGAATCA CTGGATTCAT TGCTTACGA AGCATTGCG GTTGTCCGTG AAGGTGCCAA	3000
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TGGTGACGTG CCAGAGATGC GTACAGGGGA AGGGAAAACC TTGACTGCGA CCATGCCGGT	3120
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CTCAACTAAC TCAGAAATCG GATTGACTA CCTTCGTGAC AACATGGTCG TTCGCGCCGA	3360
AAACATGGTA CAACGTCCGC TTAACTATGC CTTGGTCGAT GAGGTTGACT CTATCTTGAT	3420
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GTATCACATG GCAGACCACT ATGTAAAATC TTTGAACAAA GATGACTACA TCATCGATGT	3540
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ACTTGAAAAC CTCTATGACA TCGAAAACGT GGCTTGACT CACTTTATCG ATAACGCCCT	3660
TCGTGCCAAC TACATCATGC TTCTCGATAT TGACTATGTG GTGAGCGAAG AGCAAGAAAT	3720
CTTGATTGTC GACCAATTAA CAGGTCGTAC CATGGAAGGT CGTCGTTATT CTGATGGATT	3780
GCACCAAGCT ATTGAAGCCA AAGAAGGTGT GCCAATCCAG GATGAAACCA AGACATCTGC	3840
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GGTTGGTACA GTAGCGGTTG AAACTAGTGA CTACATTCT AAGAAATTGG TTGCAGCTGG	4140
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TGCTGGTCAA CGTGGTGCCG TTACCATCGC AACCAACATG GCGGGTCTGT GTACCGACAT	4260
CAAGCTTGGT GAAGGTGTTG GTGAACCTGG AGGACTTGT GTTATTGGTA CAGAACGTCA	4320
TGAAAGTCGT CGTATCGATA ACCAGCTTCG TGGACGTTCA GGTCGTCAAG GAGATCCAGG	4380
TGAGTCACAA TTCTACCTAT CTCTTGAAAGA TGATTTGATG AAACGTTTG GTTCTGAACG	4440
CTTGAAAGGGA ATCTTGAAAC GCTTGAACAT GTCTGAAGAG GCCATTGAGT CTCGCATGTT	4500

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GACGCGTCAG	GTTGAAGCAG	CTCAGAAACG	TGTCGAAGGA	AATAACTACG	ATACCCGTAA	4560
ACAAGTCCTT	CAATACGATG	ATGTCATGCG	TGAACAACGT	GAGATTATCT	ATGCTCAACG	4620
TTACGATGTC	ATCACTGCAG	ATCGTGACTT	GGCACCTGAA	ATTCAGTCTA	TGATCAAACG	4680
CACGATTGAA	CGTGTGTTG	ATGGTCATGC	GC GTGCCAAA	CAAGATGAAA	AACTAGAGGC	4740
AATTTTGAAAC	TTTGCTAAGT	ACAAC TTGCT	TCCTGAAGAT	TCTATTACGA	TGGAAGACTT	4800
GTCAGGCTTG	TCTGATAAGG	CCATCAAGGA	AGAGCTTTTC	CAACGTTCT	TGAAGGTTTA	4860
CGATAGTCAG	GTTTCAAAAC	TACCGGATGA	AGAAGCAGTT	AAAGAATTCC	AAAAAGTTTT	4920
GATTCTACGA	GTGGTGATA	ACAAGTGGAC	AGATCATATC	GATGCCCTG	ATCAATTGCG	4980
TAACGCGGTT	GGACTTCGTG	GCTATGCTCA	GAACAACCC	GTTGTTGAGT	ATCAGGCAGA	5040
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AGCGACTCGC	AATATCGCTG	CTCACCAAGC	AA GTATGCCA	GAAGATTGG	ATTTGAGCCA	5220
GATTGGACGC	AATGAAC TTT	GCCC ATGTGG	TTCTGGTAAG	AAATTTAAAA	ACTGTACCGG	5280
TAAAAGACAA	TAAAATGAGA	TAGTTAGAG	GCGGATATCT	TGTGAAAAGT	AAATTTTAC	5340
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GTAATCGGGC	CATGCTCATC	TGACAACGAA	GAAGCTGTCC	TTGAATACGC	TAAGCGTTG	5580
GCAGTCCTAC	AAGAAGAAGT	GGCAGATCGT	ATCTTTATGG	TTATGCGTGT	TTATACTGCC	5640
AAACCCCCTA	CCAACGGAGA	TGGCTATAAG	GGCTTGATTC	ACCAGCCTAA	CGCGACAGAA	5700
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GAAACAGGGA	TGACA ACTGC	TGATGAAATG	CTTTATCCTG	AAAACCTTCC	GCTTGTACAT	5820
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GAAGTAGAAA	CAACTGGAA	CCCGCTTTCA	CACGCTATT	TTCGTGGTGC	TCTTAATGAG	6060
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GAGAAAATGG	GCTTGGAAAA	TCCTTTATC	ATCATTGATA	CCAATCATGA	CAATTCTGGT	6180
AAGCAGTATA	TTGAACAGAT	CCGAATTGTC	CGCCAGACCT	TGATTAACCG	TGCTTGGAAAT	6240

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CAAAATGAGC	CAGAAGTATT	TGGTAAGTCT	ATCACAGACC	CTTGCCTGGG	TTGGGATAAC	6360
ACAGAAGCTC	TTGTCAGAGA	AATTTACAAA	ACGTTAGGAG	AATAAGATGG	CATTTATTGA	6420
AAAAGGTCAA	GAAATCGATA	TGGAAGTCAT	CAAGGCTGAA	ACCCAATTGT	CTGCGGAAGC	6480
CTTGAGACTC	AAGGAAAGCC	GTGACAGGG	ATTGGCAGAT	ATTATTCAG	GGGAAGATGA	6540
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TGCTCGCCGT	TTATCTGCCT	TGCAAAAGAA	GGTAGCGGAT	AAGATTTCA	TGGTCATGCG	6660
CGTGTATACT	GCTAACCTC	GTACCAATGG	AGACGGCTAT	AAAGGATTAG	TTCACCAGCC	6720
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CCGCGTGATT	ACAGAGACTG	GTTTGACAAC	GGCAGATGAG	ATGCTTTATC	CGTCAAATCT	6840
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TGATAACTCA	GGCAAGCAAT	ATATGGAGCA	GATTCGAATT	GTTCGCCAGA	CCTTGCAGAA	7260
TCGTGATTGG	AATGAGAAAA	TTAAAAAGAC	GGTCGAGGA	TTTATGATTG	AATCTTACCT	7320
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TATTTAGTC	AGGCACCATT	TTCAGGAAAG	ATTTGGCTGT	CTATCAGCCA	CACCGATCAG	7800
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CAAGGCTCTG	ATTCATCTGG	GAGCTATTG	ACAAAATATT	CAGCAAATGG	GGGCTCATAT	7920
CCCTCAAGGA	ACGCTCAAGT	TGGCTGTGGT	TAAGGCCAAT	GCTTATGGTC	ATGGAGCTGT	7980
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GCAACACTTT	GAGAGTTAC	AGAACCTTTT	TCCAATTG	AAACTGGCTC	TCTTGACAGG	9960
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TTTGATTATA	GGAACTCACG	CTCTGATACA	AGATGGGGTG	GAGTATGCTC	GTCTTGGTT	10080
GATTATTATC	GATGAGCAGC	ACCGTTTGG	TGTAGGGCAA	AGGCGTATTT	TACGGGAAAA	10140
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TCAAGTGCTA TGAGTTAAC A TTTGAAGGAA GCAAGGAAGG AAAGGTCTAT GCACGCATTG	11700
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TGGATGTGCG GGGCCAGTC A GGTTACTCAC AAGACGGCTT GCGTTCTCCT TTAGGAAATA	11880
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ACAGTCAAAT CGATTCTAA CAATGTTTA GAAACAAATG TGTACTATTC TAGTGTCAAT	12660
CTATTATATT TATAGAATT TTTGTTGCTA GATTTGTCAA ATTGTTAAA ATAATTTTT	12720
TCAGAAAGCA AAAGCCGATA CCTATCGAGT AGGGTAGTTC TTGCTATCGT CAGGCTTGTC	12780
TGTAGGTGTT AATACTTTTC AAAATCTCT TCAAACCACG TCAGCTTCGC CTTGC	12835

(2) INFORMATION FOR SEQ ID NO: 142:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5020 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 142:

GGGGATATGA AGAACAAAAG AATATTTAAA GACTTCCAAG CTTCAAAAAT GAGTTAAC	60
ATTTACACAA GCCCCTTGTT AGCCTTGTT TTTGTCTTCA TAGGAGAGTT TGTGGCTTT	120

964	
ACTTTGTATG GTATTGGCTT GTTAGCTCTC ATCGGACTTG CTAGAAATT TGGAGAGGCT	180
GGTCAAAATC TTGCAAGCTA CTTGCAGACC TTGCATCAGA GCTTGACGGA TAAAACAAGT	240
GACTTTCGTT TAATTTAGG ATTACTGGCC TTTGGTTATT CTTAACACTG TGTTCAGATG	300
GACAAGAAAA GTTGAGAAAA GACCTATTG AACCTGGGA TTTTATAGAG AGAATTTCCT	360
CAGCAATCTT CTGAAAGGAT TTAGTCTAGG CCTGGCACTT TTTCTTCTGA CCTTGTAGG	420
TTTAGTGGTC TTAGGTCAAT ATCGTTGGA ATCCATTAC TTGAATCCTT ATTCTCTTGC	480
CTTTGTCGTC TTTACTATCC CATTGGAT TTTACAGGGG ACAGCAGAAG AAGTGGTGGC	540
CCGTGCTTGG CTACTTCCTC AATTGGCCTC AAGAACCAAT CTAAAACCTAG CTATTCTTAT	600
ATCTAGCCTG TTCTTTACCC TGCTTCATAT GGGCAATTCT GGTCTCACCC CTCTATCTCT	660
AGTAAATCTC TTTTTATTG GAGTTGCCAT GGCTCTTAC CTTCTCAAAA CTGATAACAGT	720
TTGGGGTGTGTT GCAGGTATTG ATGGTGCTTG GAATTTGCT CAGGTAATC TCTTGGGAT	780
TTTAGTTAGT GGTCAACCGT CAGAACGTCT CTGATGACCT TTTTACCAACA AGGCAATCAA	840
GATTGGCTAT CAGGTGGTTC TTTGGCATA GAAGGTTCCA TTATGACAAG TCTGGTATTA	900
CTACTGCTGA TTGCTCATCT TGCTAATAAA TAAAGAAAG AAAATGAAAG GATGTGACTT	960
CGGTCCGTCC TTTCTTCGT GAAAATACTA TAAGTATGCT AAAATAGGAA TAGCACATGG	1020
AGAGAGGATT CTTATGATCA ATCACATTAC AGATAATCAA TTTAAACTAG TATCAAAATA	1080
TCAACCATCA GGAGATCAAC CCCAAGCTAT CGAGCAGTTG GTGGATAACA TTGAGGGGG	1140
AGAAAAAGCT CAGATTCTGA TGGGGCGAC TGGAACAGGG AAGACCTATA CTATGAGTCA	1200
GGTCATTTCT AAAGTCAATA AACCAACTCT GGTTATTGCC CACAATAAAA CTCTGGCTGG	1260
TCAGCTCTAT GGGGAGTTA AGGAATTTC CCCTGAAAAT GCAGTTGAGT ATTCGTATC	1320
CTACTATGAT TATTACCAGC CAGAGGCCTA TGTCCCTTCT AGCGATAACCT ATATTGAGAA	1380
GGATAGTTCT GTCAATGACG AGATTGACAA GCTTCGCCAC TCAGCTACCT CAGCCCTTT	1440
GGAGCGTAAT GATGTTATTG TCGTGGCCTC AGTCTCTTGT ATCTATGGTT TGGGTTGCC	1500
CAAGGAATAAC GCTGATAGTG TCGTTAGTCT CCGTCCTGGT CTAGAGATT CTCGTATAA	1560
ACTCTTGAAT GACTTGGTCG ATATTCAGTT TGAACGTAAT GATATTGATT TCCAACCGGG	1620
AAGATTCGC GTTCGTGGGG ATGTGGTAGA GATTTCCCA GCTTCCCGAG ATGAACATGC	1680
CTTCGAGTA GAATTTTG GAGACGAAAT TGACCGTATT CGTGAAGTTG AGGCTCTGAC	1740
AGGTCAGGTG TTGGGAGAAG TGGATCATT AGCGATTTTC CCAGCGACAC ACTTTGTGAC	1800
CAATGACGAC CACATGGAAG TTGCCATTGC AAAGATTCAAG GCCGAGTTGG AAGAACAAATT	1860
AGCTGTCTTT GAAAAGGAAG GTAAACTGCT TGAAGCCCAG CGTTTGAAAC AGCGGACAGA	1920

965

GTATGATATC GAAATGTTGC GTGAGATGGG CTATACCAAT GGGGTTGAAA ATTATTCTCG	1980
CCACATGGAT GGACGGAGCG AAGGAGAGCC TCCTTATACG CTTCTCGACT TCTTCCCAGA	2040
TGATTTCCTTG ATTATGATTG ACGAGAGTC TATGACCATA GGGCAAATCA AGGGCATGTA	2100
CAATGGAGAC CGTTCGCGTA AAGAAATGCT GTTAAATTAT GGTTTCCGTT TGCCGCTG	2160
TTTGGACAAT CGTCCTCTCC GTCGGGAGGA GTTGAGAGT CACGTTCATC AGATTGTTA	2220
CGTTTCAGCG ACACCTGGTG ACTATGAAAA TGAACAGACC GAGACAGTGA TTGAGCAAAT	2280
CATTCGTCCA ACGGGACTCT TGGATCCAGA GGTGGAAAGTC CGTCCGACTA TGGGACAGAT	2340
TGATGACCTC TTGGGTGAAA TCAATGCCCG CGTTGAAAAA AATGAGCGTA CCTTTATCAC	2400
AACTTGACC AAGAAAATGG CAGAGGATTT GACCGACTAC TTCAAGGAAA TGGGTATCAA	2460
GGTCAAGTAC ATGCACTCGG ATATCAAGAC CTTGGAACCG ACGGAGATTA TCCGTGACCT	2520
GCGCTTGGGT GTCTTGATG TCTTGTCGG ATTAAACCTG CTCCGTGAAG GAATTGACGT	2580
TCCTGAAGTG AGCCTCGTAG CTATTCTCGA TGCTGACAAG GAAGGTTTCC TTCGCAACGA	2640
ACGTGGACTC ATCCAGACCA TTGGACGTGC TGCACGTAAT AGCGAAGGTC ATGTTATCAT	2700
GTATGCGGAC ACGGTTACCC AGTCTATGCA ACGTGCTATC GATGAAACTG CCCGCCGTCG	2760
CAAATCCAG ATGGCCTATA ATGAAGAACAA TGGTATCGTT CCACAAACCA TCAAGAAAGA	2820
AATCCGTGAC TTGATTGCTG TGACCAAGGC AGTTGCTAAG GAAGAAGACA AGGAAGTCGA	2880
TATCAATAGC CTCAACAAAC AAGAGCGAA AGAACTAGTC AAAAAGCTTG GAACACAAAT	2940
GCAAGAAGCA GTTGAAGTGC TTGACTTTGA ACTAGCAGCT CAGATTGCGT ATATGATGCT	3000
GGAAGTCAAG GCCTTGGATT AGGGGAATAG TATGATTTAT TTAAGAAAGT TAAAGAAAGA	3060
AGATTTGATG TCTTTATGGG AAATGGCTTA TTCACAGCTT AATCCAGTTT GGAAACAGTA	3120
TGATGCTCCC TATTATGATG ATTATCAGTA TTTTCAAAT TTTAAAGAAT TCGAACTACA	3180
AAAATCAGAA TCCATTTAA GCAACTAAA TCGCCTGGT ATTTTGTG AGATAAAACT	3240
AGTTGGACT GTTTCGCGTT ATTGGGTATG TAAAGAAACA AGATGGATGG AATTGGGAAT	3300
TGGTATTAT GATAAAAAT TCTGGAACAC TGGTATTGGG AAAGTTGCTA TGTTGCAGTG	3360
GATAGATAGG ACGTTTCAGG ATTACTTGGG GTTGGAGCAT CTGGGTTGAA CAACTTGGTC	3420
AGGAAATATT GGTATGATGA AACTGCTGA AAAATTAAGA ATGAAAAAAG AAGCTCATAT	3480
TCCAAAAGTT CGTTATTATC AAGGTAAATA TTTTGACAGT ATTAAATATG GTATTTGAG	3540
AGAAGACTGG GAGAAAATAA ATGACGGTTA TTATCAAATC AATGGAAACT CCTGAAGAGA	3600
TAGAAGGTAA ATCCTTCGTT CACTGGAAA CGTGGAGAGA GGCTTATGAT GACCTTTGC	3660

966	
CTGCGGAATT TCAGGAGACA ATGACATTAG AAAGATGTCG ACTCTTAGT CAAAAGTATC	3720
CAGAAAATAC ATTGATTGCG ATGGATGGTG TGAAGATAGT TGGTTTATA AGTTATGGCA	3780
ACTGTCGTGA TGAGACTATT CAAGCTGGTG AAATTATTGC TTTATATGTT TTAAAAGACT	3840
ATTATGGAAA AGGAATCGCA CAAAAGTTAG TGAAAGCAGC TTTGACTGAT CTTAACATT	3900
TTTCTGAAAT TTTCTTATGG GTATTGAAAG ATAACAAGCG CGCCATTGCT TTCTATCAA	3960
AAATGGGTTT TACTTTGAT GGACAAGAAA AAATACTTGA ACTTGAAAG CCTATAAAGG	4020
AAAAACGGAT GGTATTCTAT TCTAAATAAT TCTCAAAAGT AAAAGCTAAT ATGGTACCAA	4080
GTCTGAAAAT TTAATAAATT AGAAAGCGAG TAAATTTATG TCCC GTTCCC AATTAACAAT	4140
TTTAACAAAT ATCTGTCTGA TTGAAGACCT CGAAACTCAG CGCGTGGTGA TGCACTATCG	4200
CGCCCCGTGAA AACAAATCGCT GGTCTGGTTA TGCCCTTCCT GGAGGTCATG TAGAAAATGA	4260
TGAGGCTTT TGCGAGTCTG TCATTGTGA AATCTACGAA GAAACAGGGT TGACTATCCA	4320
AAATCCTCAA CTTGTCGGCA TTAAAAATTG GCCACTAGAT ACAGGTGGGC GCTATATTGT	4380
CATTGTTAT AAGGCAGCTG AGTTCTCTGG TACCCTTCAA TCTTCAGAAG AGGGAGAAGT	4440
TTCCTGGGTG CAAAAAGACC AGATTCCAAA CTAAATCTG GCCTATGATA TGCTACCATT	4500
GATGGAAATG ATGGAAGCTC CCGACAAGTC AGAGTTTTTC TACCCTCGCC GTACAGAAGA	4560
CGATTGGGAA AAGAAAATCT TCTAGTCTTT TACTAAATAA CCTAGCTGAT CCAAGGCCTC	4620
CTCGATATAG TGGAGGTCTT GTTGTGTCTC GGCTTCAACT AGGTGATAAT GAATACCATC	4680
TGTTAACTCA GAAATTGGCT TAAAGTCAGA ACGTTCAACT TGTTCTAGAA AATGTTGCAC	4740
GTCGCGGGGA CAGTCAGTT TTAGTAAGGT TTCAATCTCT CCATAAACAG GATGATCAAT	4800
CAAGATATTT TGAACCGCAC CACCATTATC TACGATAGCA AGTAATTCTC GTCCAATTTC	4860
TTCAACTTCA TGCTTGACCT TAAATAATTG GTGATGATAA GTATTTGCAT TAGCATCTT	4920
ATAGATATAA CCACGATTGG TAGATAGAAT TGGAGATCCA TCAGCTCTTA AAATTGCAAT	4980
ATCTTGAACA ATAACCTGTC GAGTGACATG AAAGTGCTCA	5020

(2) INFORMATION FOR SEQ ID NO: 143:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4965 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 143:

AAAAAGTGGC AATCCATTGA TTGGCCACTT CATTAGAGA ATTATCGTCT CGCCCTTGAA

60

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GAAGAAGGTC	GTTAGTACT	TGAGTTACTG	CTATCGCTAG	AACTACTACT	TGAAC TGCTG	120
GAGCTGGATG	GAGTTGGTAG	ACTCCCCACA	ATACTAGACC	AAGCATTCTG	ATAATCCGCA	180
TCAC TCCGC	CAATAGCAA	GCGATAACTT	GTCGCTGGCG	CTCCTGACTT	ATTAGCCAA	240
TAGCTGGTAA	CAGTCGAACC	TGTGACCTCT	ACTTCTTTTC	CTTCAACAGA	AACCTTCTCT	300
GGTTTTTGAC	CTGTTGATTT	CAAGACTTCC	GATTCACTA	CACTAGGATC	TAAAGCAAAG	360
CGCTCGTTCC	CCCAAATGCT	TGGGAAGCT	TGCTGAATCG	CATT TACCA	ATGAGCCATG	420
TAATTAGAGT	TATTAGAATA	ACCTGCTCTA	CGTGACAATG	AATGATTATC	ATCATGCCA	480
ATCCAGCCAC	CTAGGGTTAA	TCTAGGTGTC	GAAAGCATGA	GCCACATATT	TCGCTTTGG	540
TTGGTTGTAC	CAGTCTCCC	AATCCAATCT	GCATTAGCCA	GAGTAGGATT	TAAAGAAGTC	600
AGGTTAGACT	TGAAGGGTGT	TGTCACACGA	GAGGATAGAA	CTTCTCGTAG	CAATCCCTGC	660
ATAATCGTCG	CAGTAGCTTT	TGAATAGACT	TGAACCGGTT	TATCCTGATA	CTCATACACC	720
ACTCTACCAT	CTGCTGCTTC	AATCTTGAA	ATCACATGCT	TCTGATGATA	AACTCCATTA	780
TTAGCTAAGG	TCTGATAGCC	ATTGGTATGC	TGGGCAACTG	TGACTTCAAT	ACCACCA	840
ATTGGCAAGC	TCTCAATACC	GTACTCAGGA	ATCTCGTAAC	CCATCTTTTC	CATATAACCC	900
TTGACATCAA	CACCCTTTTC	ACGGAGCATA	CGATAGGTCC	AGTAAGCAGG	GATATTCCAT	960
GAATAGTTCA	GAGCTCTCC	CAAGGTCA	ATTCTGTTC	CCTTGCTATT	AGCATAACATA	1020
ATCGGATTGC	CATTAGCAA	GTGTTGGA	TAGTTAGATA	GAATCGTTTC	ACTTCCCATC	1080
AAGCCCTGGT	CAATAGCAAT	ACCGTAGGCC	ACCAAGGGCT	TGGTAGT	AGCTGGCGAA	1140
CGTTGGTAT	CAAAGGCATG	ATTATTTGA	TTTTCTTGAT	AATTACGACC	ACCTACAAAG	1200
CCTAGAATAG	CACCTGTTG	GTTATCCATC	AAGACATTCC	CTACTTCTAC	ACGACCTGTT	1260
CCATCGTCTA	AAAGATAGCC	ATAATCAGCA	ACCGCACTTT	GCATGCCAGA	ATGAATTTC	1320
TGATCTATGG	TAGTAGTAAT	CTTATAACCA	CCATTTCAA	TTTCCTGGC	TGCCAAATCT	1380
CGATAAAACT	TCTGAGTTGC	CTCATTTTTC	AACTCCTTAG	CGGAGACATT	GTCTCTCTGA	1440
GCTAGATAGT	CATACATACG	TTCTTGAGCT	TCTGCCAAAG	TTGTAAAGTA	TAAATAGTCT	1500
CGTGAAATTC	CTGTAACCGT	GCCCCGATGGT	AAAAAGTCCT	GTTTAAGGTC	ATAATCCTTG	1560
TACTGAGAAT	ACTCGCTTT	GCTTAATGCA	CCTGTACGAT	ACATACTGTA	AAGAACTGCC	1620
TTAGCCCGTC	TTAAGCCAAT	TTCTAGGTCT	TCATCACTCT	TCAACTCCCC	AGTATTTC	1680
TAAGGAGAGT	AAGTAATGGG	ACTCTGTGGA	AGTCCTGCTA	AAAATGCTGC	TTGAGGAACA	1740
GTCAACTGAC	TGGCATCTAC	ACCGAAAATT	CCCTCAGCTG	CTTGCCGAGC	CCCTGCAATA	1800

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TTCTGTCCT TATTATTCG	GCCAAAGGGA	GCCACATTGA	GATAGGTCGT	TAAAATCTCA	1860
TCTTTATTCA	TGGCGCGTTC	CAAGGCAAGA	GCATCCACAA	TCTCTGCCG	1920
AAGGTCGGCG	CATCCCCAAC	CACCTGCTGT	TTAATTAGTT	GCTGGGTCAA	1980
CCACTAGAGG	AACCCAAAC	TACAAATTTC	CCCAAGGTCG	CACGAATCAC	2040
ACTACACCCT	TATGTTCTT	AAAGTGTCA	TCTTCTGTCG	CAATGATAGC	2100
TTTTCGAAA	TTTGCTCAGA	TGAGATAGAA	GTGCGCAACA	AATCACTCTC	2160
ATCACCGTCC	CGTCCGAATA	GGTAATCTCT	GAAATAGAAG	AGATGTCCTT	2220
ACCAATTCTT	CTGTCGAGG	CACCGAAC	TTGTCAAATA	AGGCCACTCC	2280
GCAATCCCAG	CTCCCCACAT	TCCTCCTAGA	AAACCGAGTA	CAAAGAGTAA	2340
GCTTTTATAC	TCAGTAAAAT	AGCTGGAAA	ATGACTGACT	TATCTAAGGT	2400
TTGGTACTTG	AACCTTCTT	GCCAGGTCTA	GCTGATTTT	TATTTTTTG	2460
AAAAATTCCA	GCATTTTCG	TTTTAATTCA	TTTAATTGAT	TTTGCATGGA	2520
TTATCTATTA	TACCACAAA	GGGAAATTTT	CAATAAAAATA	GCCACTTCT	2580
GCTAGGCTAT	TGCCCAAGTT	TGTGATACAA	TAGGTAGAAA	CAATAATT	2640
AAAAACACAT	GCACATT	GATGAGCTAA	AAGAGCGTGG	TTTGATATT	2700
ATGAAGAACG	TTTGC	AAAAG	AAGGTCAAGT	TTCTTATT	2760
ATCCAAC	TGACAGC	TTT	CACTAGGCC	ACCTTGT	2820
TGCAACTAGC	AGGT	CAAAA	ACCGC	GC	2880
ATCCGT	CAAAGATG	GT	AACTG	TCG	2940
TCAAGT	CCAAGG	AC	AACTAC	GG	3000
CTGTC	CAACAA	CT	ACTG	GG	3060
ATATTG	AA	TT	ACTTAC	GG	3120
TCGAA	ACAGG	TT	ACTGAG	TT	3180
TCG	AA	TT	ACT	GG	3240
ATATG	ACAG	TT	GTAC	GG	3300
CTG	AA	TT	CACT	GG	3360
TCTGG	AA	TT	CCAT	GG	3420
TGGAC	AA	TT	GTTC	GG	3480
AAGA	AA	TT	TAC	GG	3540
CTCGT	AA	TT	TAC	GG	3600

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CTGAGCAACT	CTTTGCAGGA	AACATCAAAA	ACCTTCTGT	CAAAGAGCTC	AAACAAGGAC	3660
TTCGTGGTGT	GCCCAACTAC	CAAGTACAGG	CAGACGAAAA	CAACAATATC	GTGGAACCTGC	3720
TCGTCTCATC	TGGTATAGTT	AACTCAAAAC	GCCAAGCCCG	TGAAGACGTC	CAAAACGGAG	3780
CCATCTACGT	AAACGGCGAC	CGCATCCAAG	AGCTTGACTA	TGTCTTGAGT	GACGCTGATA	3840
AGTTAGAGAA	TGAAC TGACT	GTTATCCGTC	GTGGGAAGAA	AAAATAC TTT	GTATTGACTT	3900
ACTAAACTAT	TCAACATTTA	TCTATAAACAA	AAGGAGTTAA	CCTCGAGAAA	GGTAAC TCC	3960
TTTGCTGTT	AATAACTCTC	ATCTATCTAT	TTTTAATAGA	CAGGCTACGC	AGGACAATGC	4020
GCAAGGTTGT	TAGATTATGT	AAGATAGAGA	GATTGAGG	ACTGAACCAA	TTAAATAAGC	4080
CAAAGCCAAT	CAAAC TACTA	TTTACGACAA	CGGTATCCTG	AATATTTTC	TTGATGAGTG	4140
TTTGCAAAGA	TGATGATAAC	GAATCCA ACT	CTTGGGAAGAA	ATCCAAACGA	TTATCTAAC	4200
ATAAGATATC	ACTCATCTGC	TTAGAAATAT	CTGCACTCTC	ATTCATCAC	ACACCGATAT	4260
CTGATAGAGT	TAAAGCCGCT	GAGTCATTCA	ATCCATCTCC	AACCATCAA	ATAGTGTGAC	4320
CTGCTTCTG	CAGTTCTCT	ACTAACTCAA	ATTCCCAC	AGGTTCAAG	TCTGTATAGA	4380
CCTGATCAA	GGGCAAATCT	TTGACTAATT	CCTCTGTCC	AATCAAGGTG	TCTCCTGTTG	4440
CCAGAATCAA	TTTTTCCCC	TGTGCCCTAA	GTTTATCAA	GGCTGTTTT	GCTTCTTTTC	4500
TCAAAGGAGT	ATGAATGCAG	AACATTCAA	TCAATT CATT	TTGATAAGCC	AAGAATAAGA	4560
GATTGTAGTG	ACTCTTGAC	TCTTCAATT	AAGCATT TTG	TTCTGAAC	ATATGAATCT	4620
GCTCATCCTG	CATCAAGACA	TAATTCCCAA	TAAGAACTGG	TTGGCCATCT	ATATGAGATT	4680
TGATCCCCTT	GCTT GCGATA	TATTGGAGTT	TCCCATGCAT	TTCCTCATGT	TCAATTCCCT	4740
CTATCTCAGC	TTGCTTGACG	ATGGCATTAG	CAATAGGATG	ATAAAATGTGT	TCCTCAAGAC	4800
AGGCACTGAT	TCTGAGAATA	TCTTCCAC	TATAGTCTCC	AAAAGGT AAC	ACCTTTCAA	4860
CTATAGGATA	ACTAGTTGTG	ATTGTTCTG	TCTTATCAA	CAAGAAAGTA	TCAACTTCCA	4920
GATATTCTC	CCTGTTGTGG	CCTCTGGCTG	TCATCTCTGT	GCTGG		4965

(2) INFORMATION FOR SEQ ID NO: 144:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3232 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 144:

970	
CAGGGGCGTA TTACGTGACA ATTCAATGTA GGCTGTCGCT ACTTGCGCCA AAACAAGGAT	60
TCGATAATGT CGGATGATAC TAACGATTAA ACCGAGCAGA AAGGATCCCA AAATTCCCCA	120
AACTGCAATA TGCAAGGTCA GAAAGAATGC CTTTGATAT AGTGGTAGAT ATTGTTAAC	180
AATGGATCAA TCCAAAATA GAACCTCCC TCTAGAAATA ATACAGTTAT TGTAGCACTT	240
AAAATCTTCT TTGGATAATA TCTATTTTT ATTGCCGTTA TAAGGATTT TATCATAGAC	300
ATAAAATTTC TGAAATTCC AAACAAAATA TTTAAAAGT TTTGAAAAAG AGTTAAGATA	360
TTTTTGTAA ACACAAAGTA AACGCTTAAC TATTAAGGAG GACATTTAT GTCATACAAA	420
ACAAGCAATG CAGAAGGTCA TGTAGATTT ATCAATACCT ATGATTTGGA GCCAATGGCG	480
CAACAAGTTA TTCCTAAAGC AGCATTGTC TATATCGCTA GTGGGGGGG AGATACTTTC	540
ACTTCTTCC AGTGATTTA GCGTCAGGTT CTTTTAGTT TTTAAAGATT ATCCGTGAAT	600
TTCTTGCTTA TTTATGATAA AATGGGAGTG TCGCAAAAAA TGACTCATCG TATTCAATT	660
TGAGTAAAAC TAGGAGGATC CCATGTCTAC AGAACATATG GAAGAACTAA ATGACCAGCA	720
GATCGTTCGC CGTGAAAAAA TGGCTGCGCT CGCGAACAA GGAATCGATC CTTCCGAAA	780
ACGTTTGAA CGTACTGCAA ATTACAAAGA ATTAAAAGAT AAATATGCCA ACCTCGATAA	840
AGAACAAATTA CACGATAAAA ACGAACACGC TACTATCGCA GGACGCTTGA TAACCAAACG	900
TGGTAAAGGA AAAGTTGGTT TTGCCACCT TCAAGACCGC GAAGGCCAGA TTCAGATCTA	960
CGTTCGTAAG GATGCTGTCG GTGAAGAAAA CTACGAAATC TTCAAAAAAG CAGACCTTGG	1020
TGACTTCCTT GGTGTCGAAG GTGAAGTGAT GCGTACGGAT ATGGGAGAAC TCTCTATCAA	1080
GGCACCCAC ATCACACACT TGTCTAAGGC TCTTCGTCT CTTCTGAGA AATTCCATGG	1140
TTTGACAGAC GTTGAAACAA TTTACCGTAA ACGTTACCTT GACTGATTT CTAATCGTGA	1200
AAGCTTGAA CGCTTGTCA CTCGTTCAAA AATCATCTCT GAAATCCGTC GTTACCTTGA	1260
CCAAAAAGGA TTCTTGAAAG TGGAAACACC TGTTCTTCAT AATGAAGCCG GTGGTGCTGC	1320
TGCCCGTCCA TTTATCACCC ACCACAATGC CCAAAACATT GACATGGTGC TTCTATCGC	1380
GACTGAGCTT CACTTAAACAC GCCTTATCGT GGGTGGTATG GAACGTGTCT ATGAAATTGG	1440
CCGTATCTTC CGTAACGAAG GAATGGACGC TACTCATAAC CCTGAGTTCA CTTCTATCGA	1500
AGTTTACCAA GCTTATGCAG ACTTCCAAGA CATCATGGAC TTGACTGAAG GCATTATCCA	1560
ACACGCTGCT AAATCAGTCA AAGGTGATGG CCCAGTCAAC TACCAAGGTA CTGAAATCAA	1620
GATTAACGAA CCATTTAACG GTGTTCATAT GGTGGATGCT ATCAGAGAAA TTACTGGTGT	1680
CGATTTCTGG CAAGACATGA CTTTGGAAAGA AGCTAAAGCT ATCGCTGCTG AGAAGAAAGT	1740
TCCAGTTGAG AAACACTACA CTGAGGTTGG TCACATCATC AATGCCTTCT TTGAAGAGTT	1800

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TGTTGAAGAA	ACTTTAATCC	AACCAACCTT	TGTCTATGGA	CATCCAGTAG	CTGTATCTCC	1860
ACTCGCTAACG	AAAAATCCTG	AAGACCAACG	CTTTACTGAC	CGTTTCGAGC	TCTTTATCAT	1920
GACTAAGGAG	TACGGTAATG	CCTTACTGAC	GTTGAACGAC	CCAATCGACC	AACTTAGCCG	1980
TTTTGAAGCC	CAAGCTAAAG	CCAAAGAACT	TGGTGATGAT	GAAGCGACAG	GAATCGACTA	2040
TGACTACATT	GAAGCTCTTG	AATACGGTAT	GCCACCAACA	GGTGGTTGG	GAATCGGTAT	2100
CGACCGTCTC	TGCATGCTCC	TCACTGATAC	AACAACATAC	CGTGATGTAT	TGCTCTTCCC	2160
AACAATGAAA	TAAATTCTTA	TCCTCTGGGT	CTTATCAGAG	GATTTTTGA	TTCAAAAAGA	2220
GACTGAATT	AAGGAGAAAA	TGAAGTGTAG	TATATTGAAA	TTGAAATAGT	ACACTTTGAT	2280
TTCTAAGACA	TTGTTAGAAA	TTGGTTAAA	TTCCCTAACG	AATTTGTGCA	TGTTTTATTT	2340
CATTTTACGA	TAGTACCGCTG	AAACTTTCA	AAAAGTACTA	GAAATTGACT	TGGATTCCCC	2400
AAATTGATTG	TTCAGATTCA	CTATAAATAA	AAAATTAATA	AGTGGGATAG	GAAGTTAGCG	2460
TCAACTAGGA	TAGTATCTTG	CTTAAACAGT	ATATATGGGA	TTGATATAAG	TCCATAGGTC	2520
CTATTAGAGG	ATGTTCTGGT	GTCTTATTCA	CTTGTGTTTT	ATAGTATTAG	TAGATAGAAT	2580
CAGCAAATAA	AAACCCAAAT	CATTCTACACC	TCTCTCAACT	AGATGTAACT	TACAAAACCC	2640
CTGACCTCAT	GAGCCACTTT	CTTCCTCCTC	ATGAGGTCAG	TTTTACTTTC	TGCTGTTCCA	2700
GTATCGTTT	TCCTCGCTAG	ATTTCCCTCAA	AAGGGCAGAC	TCCTCCCTTG	GTGCGTCACA	2760
CGATTTTTTC	ATCTCGACTG	TTCTTTAACG	CATCATTAAC	GACGCTTTTC	TTCTAGGTGG	2820
TTCATAAGGA	ACAGGAAGAT	TCAGGTTGAC	TTTTCTAAC	CTAGAATAAA	GTGCTGAAAA	2880
CAATTCCGAA	TAGGCATAGA	GACTAGACAA	TTTGAGGAGC	TGCTTCCGTC	CTGTTCGAAC	2940
ACATTTCCC	ACCACGTGAA	GAAAAAGATG	GCGGAAGCGT	TTGATTGTTA	AAGTTGGAA	3000
GTCACCTCCA	GCTAGATGTT	TGAGAAAAAG	ATAGAGATTG	TAGGCGATAC	AGCTCATCAT	3060
CATACGAACT	TCGTTTTGA	TTAAGGTTGA	ACTATCCGTT	TTATCGCCAA	AAAATCCCTC	3120
CTTCATCTCC	TTGATGAAAT	TCTCGGTTG	ACCACGTCCA	CGATAAAGCT	GAAACTGGTC	3180
TTGGCTTGTT	CCACTCGTCA	TATTTGTAAC	GAGAGAAATA	ACATCGTAGA	AC	3232

(2) INFORMATION FOR SEQ ID NO: 145:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 10711 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 145:

CCGGAGAAAA	TGATGAAAAG	TTCAAAACTA	TTGCCCTTG	CGGGCGTGAC	ATTATTGGCG	60
GCGACTACTT	TAGCTGCATG	CTCTGGATCA	GGTTCAAGCA	CTAAAGGTGA	GAAGACATTC	120
TCATACATTT	ATGAGACAGA	CCCTGATAAC	CTCAACTATT	TGACAACACTGC	TAAGGCTGCG	180
ACACAAATAT	TACCAAGAAC	GTGGTTGATG	GTTTGCTAGA	AAATGATCGC	TACGGGAAC	240
TTGTGCCGTC	TATGGCTGAG	GATTGGTCTG	TATCCAAGGA	TGGATTGACT	TACACTTATA	300
CTATCCGTA	GGATGCAAAA	TGGTATACTT	CTGAAGGTGA	AGAATACCGG	GCAGTCAAAG	360
CTCAAGACTT	TGTAACAGGA	TTAAAATATG	CTGCTGATAA	AAAATCAGAT	GCTCTTTACC	420
TTGTTCAAGA	ATCAATCAA	GGGTTGGATG	CCTATGTAAA	AGGGGAATC	AAAGATTTCT	480
CACAAGTAGG	AATTAAGGCT	CTGGATGAAC	AGACAGTTCA	GTACACTTG	AACAAACAG	540
AAAGCTTCTG	GAATTCTAAG	ACAACCATGG	GTGTGCTTGC	GCCAGTTAAT	GAAGAGTTT	600
TGAATTCAA	AGGAGATGAT	TTTGCCAAAG	CTACGGATCC	AACTAGTCTC	TTGTATAACG	660
GTCCTTATT	GTTGAAATCC	ATTGTGACCA	AATCCTCTGT	TGAATTGCG	AAAAATCCGA	720
ACTACTGGGA	TAAGGACAAT	GTGCATGTTG	ACAAAGTTAA	ATTGTCATTC	TGGGATGGTC	780
AAGATACCA	CAAACCTGCA	GAAAACTTA	AAGATGGTAG	CCTTACAGCA	GCTCGTCTCT	840
ATCCAACAAG	TGCAAGTTTC	GCAGAACTTG	AGAAGAGTAT	GAAGGACAAT	ATTGTCTATA	900
CTCAACAAGA	CTCTATTACG	TATCTAGTTG	GTACAAATAT	TGACCGTCAG	TCCTATAAAT	960
ACACATCTAA	GACCAGCGAC	GAACAAAAGG	CATCGACTAA	AAAGGCTCTC	TTAAACAAGG	1020
ATTCCCGTCA	GGCTATTGCC	TTTGGATTTG	ACCGTACAGC	CTATGCCCTCT	CAGTTGAATG	1080
GACAAACTGG	AGCAAGTAAA	ATCTTGCCTA	ATCTCTTGT	GCCACCAACA	TTTGTCAAG	1140
CAGATGGTAA	AAACTTGGC	GATATGGTCA	AAGAGAAATT	GGTCACTTAT	GGGGATGAAT	1200
GGAAGGATGT	TAATCTTGC	GATTCTCAGG	ATGGTCTTTA	CAATCCAGAA	AAAGCCAAGG	1260
CTGAATTG	TAAAGCTAAA	TCAGCCTTAC	AAGCAGAAGG	AGTCCAATTC	CCAATTCTT	1320
TGGATATGCC	AGTTGACCAA	ACAGCAACTA	CAAAGTTCA	GCGCGTCCAA	TCTATGAAAC	1380
AATCCTTGG	AGCAACTTTA	GGAGCTGATA	ATGTCATTAT	TGATATTCAA	CAACTACAAA	1440
AAGACGAAGT	AAACAATATT	ACATATTTG	CTGAAAATGC	TGCTGGCGAA	GACTGGGATT	1500
TATCAGATAA	TGTCGGTTGG	GGTCCAGACT	TTGCCGATCC	ATCAACCTAC	CTTGATATT	1560
TCAAACCTTC	TGTAGGAGAA	AGTACTAAAA	CATATTTAGG	GTTTGACTCA	GGGGAAAGATA	1620
ATGTAGCTGC	AAAAAAAGTA	GGTCTATATG	ACTACGAAAA	ATTGGTTACT	GAGGCTGGTG	1680
ATGAGACTAC	AGATGTTGCT	AAACGCTATG	ATAAATACGC	TGCAGCCCAA	GCTTGGTTGA	1740

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CAGATAGTGC	TTTGATTATT	CCAACTACAT	CTCGTACAGG	GCGTCCAATC	TTGTCTAAGA	1800
TGGTACCATT	TACAATACCA	TTTCATTGT	CAGGAAATAA	AGGTACAAGT	GAACCAGTCT	1860
TGTATAAAATA	CTTGGAACTT	CAAGACAAGG	CAGTCACTGT	AGATGAATAC	CAAAAAGCTC	1920
AGGAAAAATG	GATGAAAGAA	AAAGAAGAGT	CTAATAAAA	GGCTCAAGAA	GATCTCGCAA	1980
AACATGTGAA	ATAACTGTTG	CAAAATATAA	GAAAGGATT	AGTATTTCCC	TTGAATGCTG	2040
AATCCTTTT	TACATTGTA	AAGAAAGATT	CTAAAATGTA	CGGACCCCCA	AAAGTTGGAG	2100
CCTCTTTTG	TCAGAATAGA	GAAAATTTT	GTAAATTTA	CTTGTTCCT	ATTGCTTCT	2160
CAGCTATTAT	TTGTTATATT	AAAAGTATAA	TTATTTTTA	TTTATCAGAG	TTAACGATTG	2220
CACTTTCAGA	GGAAGGAGTA	TTTTTAAAAA	AGAAAATGTA	AACGTTGCT	CAAAAATGAA	2280
AGGATTTAGA	AGTTTATGAA	TAAAGGATTA	TTTGAAAAC	GTTGAAATA	TAGTATTCCG	2340
AAATTTTCAT	TAGGTGTTGC	TTCTGTTATG	ATTGGAGCTG	CATTCTTGTG	GACAAGTCCG	2400
GTTCTTGCA	ATAGCGTGCA	GTCTGGTTCC	ACGGCGAACT	TACAGCTGA	TTTAGCTACT	2460
GCTCTTGCAA	CAGCAAAAGA	GAATGATGGG	CGTGATTTG	AAGCGCCTAA	GGTGGGAGAA	2520
GACCAAGGTT	CTCCAGAACT	TACAGATGGA	CCTAAGACAG	AAGAAGAACT	ATTAGCACTT	2580
GAAAAGAAA	AACCGGCTGA	AGAAAAACCA	AAAGAGGATA	AACCTGCAGC	TGCTAACCT	2640
GAAACACCTA	AGACGGTAAC	CCCTGAATGG	CAAACGGTAG	CGAATAAAGA	GCAACAGGGA	2700
ACAGTCACTA	TCCGAGAAGA	AAAAGGTGTC	CGCTACAACC	AACTATCCTC	AACTGCTCAA	2760
AATGATAACG	CAGGCAAACC	AGCCCTGTT	GAAAAGAAGG	GCTTGACCGT	TGATGCCAAT	2820
GGAAATGCAA	CTGTTGATTT	AACCTTCAAA	GATGATTCTG	AAAAGGGCAA	ATCACGCTTT	2880
GGTGTCTTT	TGAAATTTAA	AGATACCAAG	AATAATGTT	TTGTCGGTTA	TGACAAGGAT	2940
GGCTGGTTCT	GGGAGTATAA	ATCTCCAACA	ACTAGCACTT	GGTATAGAGG	TAGTCGTGTT	3000
GCTGCTCCTG	AAACAGGATC	AACAAACCGT	CTCTCTATCA	CTCTCAAGTC	AGACGGTCAG	3060
CTAAATGCCA	GCAATAATGA	TGTCAATCTC	TTTGACACAG	TGACTCTACC	AGCTGCGGTC	3120
AATGACCATC	TTAAAATGA	GAAGAAGATT	CTTCTCAAGG	CGGGCTCTTA	TGACGATGAG	3180
CGAACAGTTG	TTAGCGTTAA	AACGGATAAC	CAAGAGGGGG	TAAAAACAGA	GGATACCCCT	3240
GCTGAAAAAG	AAACAGGTCC	TGAAGTTGAT	GATAGCAAGG	TGACTTATGA	CACGATTCAG	3300
TCTAAGGTCC	TCAAAGCAGT	GATTGACCAA	GCCTCCCTC	GTGTCAAGGA	ATACAGCTTG	3360
AACGGGCATA	CTTGCCAGG	ACAGGTGCAA	CAGTTCAACC	AAGTCTTAT	CAATAACCAC	3420
CGAACATCACC	CTGAAGTCAC	TTATAAGAAA	ATCAATGAGA	CAACAGCAGA	GTACTTGATG	3480

974	
AAGCTTCGCG ATGATGCTCA CTTAATCAAT GCGGAAATGA CAGTACGCC TT GCAAGTTGTA	3540
GACAATCAAT TGCACTTGA TGTGACTAAG ATTGTCAACC ACAATCAAGT CACTCCAGGT	3600
CAAAAAGATTG ATGACGAAAG CAAACTACTT TCTTCTATT A GTTCCCTCGG CAATGCTTTA	3660
GTCTCTGTTT CTAGTAATCA AACTGGTGCT AAGTTTGATG GGGCAACC CATCAACAAAT	3720
ACGCATGTCA CGGGAGATGA TCATATCGAT GTAACCAATC CAATGAAGGA TTTGGCTAAG	3780
GGTTACATGT ATGGATTGT TTCTACAGAT AAGCTTGCTG CTGGTGTGTTG GAGTAACCT	3840
CAAAACAGCT ATGGTGGTGG TTCGAATGAC TGGACTCGTT TGACAGCTTA TAAAGAAACA	3900
GTCGGAAATG CCAACTATGT AGGAATCCAC AGCTCTGAAT GGCAATGGGA AAAAGCTTAT	3960
AAGGGCATTG TTTTCCCAGA ATACACGAAG GAACTTCAA GTGCTAAGGT TGTTATCACT	4020
GAAGATGCCA ATGCAGACAA GAACGTTGAT TGGCAAGATG GTGCCATTGC TTATCGTAGC	4080
ATTATGAACA ATCCTCAAGG TTGGGAAAAA GTTAAGGATA TCACAGCTTA CCGTATCGCG	4140
ATGAACCTTG GTTCTCAAGC ACAAAACCCA TTCTTATGA CCTTGGATGG TATCAAGAAA	4200
ATCAATCTCC ATACAGATGG TCTTGGCAA GGTGTTCTCC TTAAAGGATA TGGTAGCGAA	4260
GGCCATGACT CTGGTCACTT GAACTATGCT GATATTGGTA ACCGTATCGG TGGTGTGAA	4320
GACTTCAAGA CCCTAATTGA GAAGGCTAAG AAATATGGAG CTCATCTAGG TATCCACGTT	4380
AACGCTTCAG AAACTTATCC TGAGTCTAAA TACTTCAATG AAAAATTCT CCGTAAGAAT	4440
CCAGATGGAA GCTATAGCTA TGGTTGGAAC TGGCTAGATC AAGGTATCAA CATTGATGCT	4500
GCCTATGACC TAGCTCATGG TCGTTTGGCA CGTTGGGAAG ATTTGAAGAA AAAACTTGGT	4560
GACGGTCTCG ACTTTATCTA TGTGGACGTT TGGGTAATG GTCAATCAGG TGATAACGGT	4620
GCCTGGCTA CCCACGTTCT TGCTAAAGAA ATTAACAAAC AAGGCTGGCG CTTTGCATC	4680
GAGTGGGCC ATGGTGGTGA GTACGACTCT ACCTTCCATC ACTGGGCAGC TGACTTGACC	4740
TACGGTGGCT ACACCAATAA AGGTATCAAC AGTGCATCA CCCGCTTTAT CCGTAACCAC	4800
CAAAAAGATG CTTGGTAGG GGACTACAGA AGTTATGGTG GTGCAGCAA CTATCCACTG	4860
CTAGGTGGCT ACAGCATGAA AGACTTTGAA GGCTGGCAGG GAAGAAGTGA CTACAATGGC	4920
TATGTAACCA ACTTATTGTC CCATGACGTC ATGACTAAGT ACTTCAAACA CTTCACTGTA	4980
AGTAAATGGG AAAATGGTAC ACCGGTACT ATGACCGATA ACGGTAGCAC CTATAATGG	5040
ACTCCAGAAA TGCAGTGGA ATTGGTAGAT GCTGACAATA ATAAAGTAGT TGTAACCTCGT	5100
AAGTCAAATG ATGTCAATAG TCCACAATAT CGCGAACGTA CAGTAACGCT CAACGGACGT	5160
GTCATCCAAG ATGGTTCAAGC TTACTTGACT CCTTGGAACT GGGATGCAAA TGGTAAGAAA	5220
CTTTCTACTG ATAAGGAAAA GATGTAACAC TTCAATACGC AGGCCGGTGC AACAACTTGG	5280

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ACCCTTCCAAGCGATTGGCAAGAGAGCAAG	GTTTACCTTACAAGCTAAC	TGACCAAGGT	5340
AAGACAGAAGAGCAAGAACTGTAAGAAGATGGTAAAAA	TTACCCCTAGA	TCTTCTAGCA	5400
AATCAACCATACTGTCATACTGTCAGAACTAATC	CTGAAATGTC	ATGGAGTGAA	5460
GGCATGCACATCTATGACCAAGGATTTAAT	AGCGGTACCT	TGAAACATTG	5520
GGCGATGCTTCTAAGGCAGAAATTGTCAG	TCTCAAGGGG	CAAACGATAT	5580
CAAGGAAACAAAGAAAAGTAGTCTCACT	CAGAAATTAA	CTGGCTTGAA	5640
AAGTATGCCGTTTATGTTGGTGTAGATAAC	CGTAGTAATG	CCAAGGCAAG	5700
AATACTGGTGAAAAAGAAGTGAECTACTTAT	ACCAATAAGT	CTCTCGCGCT	5760
AAGGCCTACGCCCACAATACACGTCGTGAC	AATGCTACAG	TTGACGATAC	5820
CAAACATGTACGCCCTCTTACAACGGA	GCGGACGTCT	CAAATGTTAC	5880
AGTCGTGAAGCTGGTGTACAGCAACTTAC	TTTGATGAAA	TTCTGTACCTT	5940
TCAAGCATGTACGGAGACAGCATGATACA	GGTAAAGGCA	CCTTCAAGCA	6000
AATGTTGCTCAGGGTATCTTCCCATTGTA	GTGGGTGGTG	TCGAAGGTGT	6060
CGCACTCACTTGTCTGAAACACAATCCA	TATACACAAAC	GTGGTTGGAA	6120
GTCGATGATGTTATCGAAGGAAATTGGTCA	CTCAAGACAA	ATGGACTAGT	6180
AACTGGTTTACCAAACCATCCCACAAAC	TTCCGTTTTG	AAGCAGGTAA	6240
GTAACCTTTGAAATACGAAGCAGGATCAGAC	AATACCTATG	CTTTTGTAGT	6300
GAATTCCAGTCAGGTCGTCGTTGACTCAA	GCAAGCAACT	TGGAAATGCA	6360
AATACTGGACAGATTCTAA	GAAAGCCAAG	AAGGCAACCT	6420
ACAGGCGATACTGGTAGG	TATCTACTCA	ACTGGAAATG	6480
TCTGGTGGAAATGCCAACTTCCGTGGTTAT	AACGACTTCA	TGATGGATAA	6540
GAAGAAATTACCTAACACAGGTAAGATGTTG	ACAGAAAATG	CTCTGAAGAA	6600
ACGGTTGCCATGACTAACTCACCAAAAGAG	TCTATGGATG	CTTGAAAGA	6660
AACCTCAGTCAGGCCATGATGATCAGT	GTGGAAGAAG	CGCGTGCAGA	6720
ATTGAAGCTTGAAGAATGCTTGTTCAAG	AAGAAGACGG	CTTTGGTAGC	6780
GCAAGTCTTACAGCTCTGC	TCAGGCTCAA	GAAGGTCTTG	6840
GTGTCTAGTC	TATGGCATACTCTTGGAAAT	GGTGGAGATG	6900
GTCTTGAAAGAACCAACTGA	AATCACAGGA	CTTCGCTATG	6960
AATGGTAACCTGCGAGATGT	GAAACTTGTT	TGACAGATG	7020

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TTTACTGCAA	CTGATTGGCC	AAATAACAAC	AAACCAAAAG	ATATTGACTT	TGGTAAGACA	7080
ATCAAGGCTA	AGAAAATTGT	CCTTAAGTGGT	ACCAAGACAT	ACGGAGATGG	TGGAGATAAA	7140
TACCAATCTG	CAGCGGAAC	TATCTTTACT	CGTCCACAGG	TAGCAGAAC	ACCTCTTGAC	7200
TTGTCAGGCT	ATGAAGCAGC	TTGGTTAACG	GCTCAGAAAT	TAACAGACAA	AGACAATCAA	7260
GAGGAAGTAG	CTAGCGTTCA	GGCAAGCATG	AAATATGCAG	CGGATAACCA	TCTCTTGACG	7320
GAAAGAATGG	TGGAATACTT	TGCAGATTAT	CTCAACCAAT	AAAAGATTC	TGCTACGAAA	7380
CCAGATGCTC	CAACTGTAGA	GAAACCTGAG	TTTAAACTTA	GATCTTAGC	TTCCGAGCAA	7440
GGTAAGACGC	CAGATTATAA	GCAAGAAATA	GCTAGACCAG	AAACACCTGA	ACAAATCTTG	7500
CCAGCAACAG	GTGAGAGTCA	ATCTGACACA	GCCCTCATCC	TAGCAAGTGT	TAGTCTAGCC	7560
CTATCTGCTC	TCTTTGTAGT	AAAAACGAAG	AAAGACTAGT	ATTTAGTAAA	ACCTCTAAC	7620
AAGATTACGG	AAGCAGTCTC	TATCTTTCC	AATGAGGTTT	ATAGTACAGA	AAAAGCCTGA	7680
GAAGATGTCT	TCTCAGGCTT	TTGTTAACGA	CATAAATACA	ATAGTGCTAT	GACAAAATCA	7740
CCCAGAAAAA	TCTGGGTGAT	AAATGTTATG	GTTGTGCTGG	TTGAGGATTC	TGATTTGTT	7800
GATCAGGGGT	TGTATTGAT	TGTTGGTAT	TATTGTTAGG	ATTGGTAGTC	GTACTATTAT	7860
TTGTGCTTGG	AGTGGGTGAG	CTAGACTGTG	AAAGTTGAAC	ATCTGATGAT	GAGCTTGAAC	7920
TTTCAGTTGA	TGGGGTTGT	TGTGGAGCAG	GTGAGTTCCA	CGTAGAACGA	GCACCATT	7980
TAAATACGAA	TTCTCCATT	CTGTAGAGCC	CCTCTGGTAT	ATTCCAATCT	TCTGGATTGC	8040
TTCCTTCAGA	CAGGTAGGTC	ATCATAGAGC	GGTAAACTTT	GGCAGCGACC	GTAAGGCCAT	8100
TGCCTACAAG	TGGTGTCAAG	CGGTTAGAAT	AGCCTGTCCA	TACAGCCATT	GAATATTAC	8160
GCGTATAGCC	AGCAAATAGT	TCATCAGGTG	CTACAAATTG	AGAGGTCTTG	ATGTGGTTT	8220
CAATTTCTC	GTCTGTATAG	TTAGAGGTT	CTGTTTAC	AGCCTGAGGG	AGCCAAGCAA	8280
GATAGGCATT	TCGTCCAGTT	CCATAAGTCA	AGACTGTTT	CATCATGTCG	GTCATCATAT	8340
AGGCTGTCGT	TTCCTTCATG	GCACGAGTTC	CGACATTAGA	GAACCTTTT	TCACTCCCAT	8400
CACTAAAGAC	GACTTTATGG	ATATACATTG	GTTTATAGTA	AGTTCACCA	TTTGCAAAGG	8460
CAGCGTAAGC	AGCAGCCATC	TTTCACTAC	TTGCTCCATA	TTTTTGTC	GATTCGGTTG	8520
TGTTACTTGA	AATGGCATT	GAGTAGTGAA	TACTTGGTA	GTCGATTCC	AGACCATT	8580
GGAAAGTCTT	GGCGCGGTTG	AGTCCGACCT	TGTTTAGAGT	TTCCACGGCT	GGGACGTTTC	8640
GCGATTGTTG	CAGGGCGTAT	TGCAAGGTGA	TGTTGCCAAA	GTAGCCCCTA	TCCCAGTTAT	8700
AAACAGGAGT	ATTGTCCTCA	GGGTAGTTAT	AGGGCTCATC	GTGAACGATA	GTAGCAGTTG	8760
AATCGTAGAC	ACCGTACTCC	AAGGCAGGAG	CATAGTCTGT	GATCGGTTTC	ATAGTTGATC	8820

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CCAGTCGCG	GTTTGTTC	ACTGCTGGT	TAATTCCGAA	GGAAACATTA	CTTGACTGAT	8880
GGCGTGCTCC	TAGCTGGCA	ATGACTTTAC	CGTTAGAAC	ATCAACAATG	GTAGAACGGA	8940
CTTGCAATT	ATCGTCTGGA	TAGGCAACGT	ATTCGCTGT	ATTGTAAATA	TCCCACAGAT	9000
GTTTTGAGC	TTCTTGGTCT	ACATTGTGT	AGACATCCAT	CCCAGTTGTG	AGTAGGTTAT	9060
AGCCTGTTTC	TTCTTCAACT	TGATTGATGA	CTTCCTTGAG	GTAATTATCC	ATGTAAGCAG	9120
GGTAATTACT	TGCTGATTG	AGACTTTGTA	GTCCCATCGA	AATTGGTGT	TTGACTGCTT	9180
TCTCATACTG	TTCAGCAGAG	ATGTAGCCTT	GATTTTCAT	TTCAGATAAG	ACCAAGTTTC	9240
GGCGGTCTTG	GGCTGCTTCT	GGATGTGAAT	AGGGGTCTA	TTGGTTGGT	GCCTGAGGCA	9300
TTCCAGGCCAG	CAAGGCTAAC	TGAGGTAAAC	TTAAATTATT	GAGGTCTTTA	CCATAGTAGT	9360
TTTGAGCTGC	TGTCTGCATT	CCATAGTTCC	CATTAGACAT	GTAGACCTTA	TTTATATAGT	9420
AGGTCAAGAT	TTCTTGGTTG	GTTGCTTTT	GTTCTAACTG	AATCGCTAAC	CAAGCTTCCT	9480
GAGCCTTACG	AGAAATAGTC	TGGTCGGAAG	TCGAAGTTGA	AAAGTAAGTC	AACTTAATCA	9540
ACTGTTGGGT	GAGAGTTGAT	CCACCTTGG	GGGAATTGCT	TTGCAGATTG	CGCAAGAAAG	9600
CTCCCAGGAT	ACGGATGGTA	TCAATCCCC	TGTGGTCGAA	GAAGCGATGG	TCTTCGATAG	9660
AAACGATTGC	CTTAACCAA	TCTGTGGAA	TATCATTAGC	TTGGGCATTG	ACGCAGCGTT	9720
CAGAACCCAA	GTCAGCAATG	AGTTGATTTT	TATTGTCGTA	GATTTACTA	GAAGTTGTTG	9780
CAACTAGTT	ACTCTCGGAT	AGGCTAGGAG	CCTTGCTAAC	GTAGTAGAAA	AAAACCTCCTC	9840
CGCCTAAGAC	AATGGCTGCG	ATAACCAAGC	TTAAGAAGCT	AATGCTCAGA	TACTTGATTA	9900
GGCGCAGAAT	CGTTGGTTG	TTCATCTTGT	TTTACCACCT	AATAAATGTT	CTTTGATAAC	9960
ATTGAGATAA	GGAATTGAG	GGAAGGCACC	AGCCTTGATT	TCATATCCAT	ATTCTCGAAT	10020
ATATTCAAGT	GGCATTGATT	TTTGCCCTT	ATCTTGATGA	TAGAAGCGAA	TCAAATCGAA	10080
TGCCGGCAAT	AAGTAGGTTT	CTTGCTGAGA	AGAAAAGTGA	AGAAGGACAA	AGCAGATTCC	10140
TTGTTGGGCA	AGGACTTGTT	CCATATGCTG	AATCTGATGT	GGATGAAAAT	TTTCATCGG	10200
AATCGCACGT	TTTTGTTTG	TTTCCCTGAC	TCACAAAGTCG	ATGTAATATC	CATTATAAAC	10260
GCCAGAATAG	TCCGTCGTTG	AAGCTTGTG	AAAATAGGCT	TCAACAATCT	TGGCACGACT	10320
TCGTTGTGGA	TAGTCCACTT	GTACGATTG	AATAGGAGTT	GGTTTCTTAT	GTATAACAGC	10380
CAAGCCCTGA	GACAAATAGT	AGTCGTTGGT	AGCATTGATC	ATCTTTCAA	AGGGTACCGA	10440
GCTCGAATT	GTAATCATGT	CATAGCTGTT	TCCTGTGTGA	AATTGTTATC	CGCTCACAAT	10500
TCCACACAAAC	ATACGAGCCG	GAAGCATAAA	GTGTAAAGCC	TGGGGTGCCT	AATGAGTGAG	10560

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CTAACTCACA TTAATTGCGT TGCGCTCACT GCCCGCTTTTC CAGTCGGGAA ACCTGTCGTG	10620
CCAGCTGCAT TAATGAATCG GCCAACGCGC GGGGAGAGGC GGTTTGCATA TTGGCGCTC	10680
TTCCGCTTCC TCGCTCACTG ACTCGCTGCG C	10711

(2) INFORMATION FOR SEQ ID NO: 146:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11887 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 146:

TACATTTCATT CCATCGGCTA CTCCATAATA CTTAGATAAA ACCATAGCTG AAGTCGAATA	60
CGGATACTGT AAAGTATTAT CAATTTAAT CAAATCATCA TTACCGATAA TACTTCTGAT	120
TGCTTTGGT AGTATGAACC ATACGTTGGT GAAATCTCAG ATAATGAAGA ATCATTAGAC	180
TCTGGACCTT TTTCTAGTGT CTCACCTTAC TCATATTCTT CACCCTTACT AGAAATAACA	240
CTCAAAGCAG ATACTGTCGA TAACTGGCTA GCCAATAAG TACTCGCAAT ATTGAAATA	300
CCCAATTCTT TATAAACAGT TTTCTTCATT ATTGTATCCT CCTAATGTAA TTATAGCGTA	360
CTATTCTAAA TTTCTTAATC TACTATAGAA TCAAGAAATC TACCACCTTC TTTAAATACC	420
CTCCATTATC ACATAAACAG GTAAACTTTT CAATTATGA CTGCGCTTTT CAATCACGCT	480
AGAGGTACTT GCTTGCTTCT TTGATACTAA GTTCAGCCAT TCTTCCCTTG TTTTCTCAA	540
TAAAGCATGT TACCCAAGTG GGATTCGTTT TGGAGTAGTC TCGCAGACTC CAGCCAATGG	600
CTTTATTGAT AAAAATTCT GTTGGTTCA AGTTATGAAG GAGAATCTT TCCATTAATT	660
GAGTATTGGT CTTCTCTTTT CTTAACAACT GGTGGTCAAT AGCGACACGT CTCAGCCAGA	720
TATTATCTGA TAGGCTCCAT TTTATACTCA ATGAAAATCA AAGAGCAAAC TAGGAAGCTA	780
GCCGCAGTTG CTCAAAACAC TGTTTGAGG TTGCAGATAG AGCTGACGTG GTTTGAAGAG	840
ATTTTCGAAG AGTATTAAGA TTATTCTTC TAGTTCAGGG TGTTCATACA CCAAACCTCC	900
TACTACTCGA TCTAGGATAT CTACCGTGTG CCACAAAGGAT TTTGTCACGA CTAAC TGCTC	960
TAGCTTAGGC AAATCGTTT CCTTTAGATA AGACTGCATT GCTTCAAAT AGTTAGCAGC	1020
CACATATTGG TATTTCTAG GATCCTTTTC CCAGCAAGTG TCTGAAAAAT CCCAATCGAT	1080
AATCTTGTT TTTTCGCTT CTGGAAAATA TTTTATAGAG TTTATTTCTT TCAGGCACCG	1140
CAATACCTAG AAAAGAAAAT TGATGGCGCA TATAGGCTTC CATGGACCTT GCTTTTTAG	1200
AGTCTTTGCG TGCTTCTAGC TCCTCAAGTA AATCTGCTAA ACTCATCTAA AACTCCTCTT	1260

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GCCCCACCAA ATGGTGTGA AAGGCATAGA CAGCCGCCTG GGTACGATCG CTGACTTCAA	1320
GTTTGGCAAG AATATTGGAC ACgtgggtct TGACCGTCTT GAGAGAGATA AAGAGGTCAT	1380
CTGCGATGCG CTGATTTCTG TAGCCCTGG CGATGAGTTG GAGAACATCT CGCTCACGCG	1440
CAGTCAAATTC TTCATGAAGT TCCATATGAT TGCGGTGGTA TTCAACCTTC TTGCTAACCT	1500
CTTGCTCAAT GGCCAGCTCG CCAGCAGCTA CCTTACTGAC GGCAATGAAGC AATTCACTG	1560
CACTAGAAGT CTTGAGCATA TAGCCTTGG CACCAGCATC TAAGACTGGC ATGATTTTT	1620
CATTGTCCAA ATAAGAGGTC ACAATCAAAA TCTTGGCTTC AGGCCATTCT TTAAGGATTG	1680
CTAAGGTCGC GTCAATCCC TTCATCTCAG GCATGACAAT ATCCATGACA ATGACATCTG	1740
GACGCAGTTC CAAGGCCAAG TCAATCCCTT GAGACCCGTT GGACGCCTCA CCCACAACCT	1800
CTACATCGTC TTGGAGGTCA AAGTAGCTTT TCAAGCCCAA TCGGACCATT TCATGGTCAT	1860
CTACTAGTAA AATTTCTATC TTTACTCCTT TATCATTCCCT TATCTAACAG GGGAAATACGG	1920
ATATCAACCG CCAGCCCTTG CTTGGGAGCT GTCAAGAGTT GAACTGTTCC AGCCATATCT	1980
TCAACCCGCT CCTTGATATT TCGCAGTCCA TAACTCAAGT CGTCTAACGCT CCCTAACGG	2040
AAACCAATCC CATTGTCCAC CACCTTCAGT TGCAATTCAA CATCTGTCTG ATAGAGGTAG	2100
ACATCTAGGC AAGATGCCTG GGCATGGCGG AGGGTATTGC TAATCAACTC TTGCAGGATA	2160
CGGAAGATAT GCTCCTCGAT TTTCTTAGGC AATTTCGTCA TATTCTGCTT GAGACTAACCC	2220
CTAAGATCAC TCTTGTCCCTC AAGCTCTTTT AAAAGAATTG GAATCCCTTC TATCAAGCTC	2280
TTCTGCTCCA GTTCAACTGG TCGCAAATGC AAGAGCAAAA CCCGCAAATC CTTCTGGCT	2340
GTTTCTAAAA TAGCTGTGAC ACTCTGCAAC TGGGTCTGCA TCTTTCTCT ATCCAATTTC	2400
AAAGCCTGCT GACTGATACC CGATAAAATC ATGTGGGCCG CAAACAACTC CTGACTGACT	2460
GTATCGTGCA AATCCCGAGC AATTGCTTCT CGTTCTTCT CGATGATTTC CTCTTCTGAA	2520
GCAAGGCTCT GATTTTCAGC TTTTGAGAAGA GCCTCTGTCA AAAGGTTAAG TTTACCTGAT	2580
AAGGACTGTGA AACTGGCATC CAAATCTGGA TCTGCAACCT GAACCACTTC TTGCCCTGCT	2640
AATAAACGCT TGAGATTAGC CTGCATTTTT CTTAGAGAAA GCTCTTCGAT CCCTGCCAA	2700
AACAGGGCTA AGAGACAGGT CATGGACATG CTGAAAACCA ACAATAAAA GACAAATTTT	2760
TCTGTTTTTT CGACATCGTG CAAAAAGATA GACCAAGTCAA AATCAAGTAT TTCCAGCAAG	2820
CTGTGGGAGA AAAAAAAGAC AAATAGGAAG GAGGTGAGAG CAATAATGAC ATAGGCTTGT	2880
TTTTTCATCC TCTAACACC TCCACATCAC CAATCATAGT GGTCAAGAAA ATCTTGACAC	2940
TCTTGTTACT CTTGAGATAG TCTTTGTTT CTTGATGATA GTGTTCATCG CGGAGGGCTC	3000

980						
GCTTGGGCTG	GTTGAAAAAA	ATCAAATCCC	CATAGAGACA	GTAAACGCTG	AGACTGACTT	3060
CCACATCTAC	AGGTACGATG	ATTTGGTCG	TTCCTTACCAT	CTTTCTGAGG	ATAATGACAT	3120
TGTCATGATT	GGTTAAGATG	ACCCCTCTCCA	GATGAATAGT	GTCCTTGCCC	ATGAAGCGAA	3180
AGAGATTGAT	ATCATCGAA	TGGCAAGTCT	GGTAGCTTGA	AAAATGATGA	AGATTTCCAA	3240
ACCAACGATT	TTTCTCCTTC	TTAACCGTCA	CGACCTCTTC	AAAAACCAAA	TTGGTCTGCT	3300
CTTTTCCTG	GTTCATCATC	GGGTAAAGAA	GAAAGAGGCT	ATAGATAACC	GCAACAAAAA	3360
TAGCTAGAAT	CACAAAAGGA	TTGAGCATAA	CGATGAAAAA	GAAGAGAATG	GTTGCCGCTA	3420
CTAAAAGAAG	ATTATTTCCC	TCTTACCAAG	TGTAGTAGCG	AATCAAAGC	AAAAAGAGGA	3480
ATAGTATCAG	CAGAAAACGC	GAAAAATGCT	CTGATACCAT	CAAATCAGA	GCTCCTGTCA	3540
GAAGACAGGC	TTCGATAAAAT	AAAAAGATTT	TAAATTTCT	CATAGGTTCA	TCCTCTCCCT	3600
TCTATTTAT	CACAATTCAA	AAAAGTCACC	TCAGTCTGAG	GATGGAAAAA	AGGCGCTGGT	3660
TACGCCTTT	TCATCTGATC	CTTGCTTCT	TTAATTTTC	CATAAAGAAG	ATAGTCTACT	3720
TTTTGTAGAT	CTGCTATGGT	GGCACAGTTA	AGGAAACACA	TAATCAAGCG	TAGATCTGCT	3780
TTCCAGCCTT	GGACAATGCC	AATCACTTCT	TCAACTGTGT	AGGTTTCAAC	CAATTCCAGA	3840
ACGGTTCGTG	ACAATCCAC	AGCCTTAGCA	CCAAAAAACCA	AGCACTTAAT	CATATCCAGC	3900
GGATTCCGAA	CCCCTCCACT	AACCAAGAGT	TCGACCTTAT	CTTCCATTC	TTGGGCATTG	3960
AGAAGGGCCT	GCATGGTAGA	CTGACCCCCAT	TGATTGAGGT	AATCACGCTG	GCCACTACGA	4020
CGGTTTTCGA	TATAGGCAAA	GCTGGTCCA	CCACGACCCG	ATAGGTCCAC	TGTACGAACA	4080
CCGAATTCA	AGGCTTTTC	GATTGTCTTG	GCATCCATTC	CAAAGCCCAC	TTCCCTTGAGG	4140
ACAATAGGAA	CGGGAATTG	CTTGCTATAA	TCTGCTAGAT	GCGATTGCCA	GCTTCTAAC	4200
TTCCTTCTC	CCTCGGGCAT	GAGTAATTCC	TGCATGACAT	TGACATGCAC	TTGCAATAGA	4260
ACAGGATTCA	TCTCTTCTAC	AGTCTGAAGT	CCTAACTCGA	CAGGCTTGTGTC	CAATCCAATA	4320
TTGGTTCCAA	GGAGGAGATT	GGGATGACTA	GACTTGACAG	AAAAAGAAC	ATCCGTTGGA	4380
TTTTTGAGGG	CTGCGCTATA	AGAACCCGTT	ACAAATAAAA	TACCACAGGA	TTCCGCCACC	4440
TGAGCCAGCT	TTTGATTGAT	TTCTCTTCCC	TTATTACTTC	CACCAGTCAT	GGCATTGATA	4500
TAAAAAAGGAA	AGTCCCAC	TCGACCAGCA	AACTCTGTG	AAAGATCGAT	TTCATCCAGA	4560
TTGTAAAGAG	GCAAGGAAGA	ATGAATCAGC	TCCACCTCAT	CAAAGCTATT	ATAGGAAC	4620
TTCTGCTCAA	GGGCATAGAG	GATATGCTCG	TCCTTACGAT	TTGTCGTCA	GTCCTATCCT	4680
TTCTTGATAT	AAGAGCTCAA	TCCCCAGATC	GGCCCAACGA	TTTTTTAAGG	TTTTGGTTGA	4740
TTGCGCATCA	AAACTCAGGG	CGATGCCACA	GTCACCACCA	CCAGCACCAC	TACTCTTGGC	4800

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AACGGTCTGC AAATCTTGAC TGGCTCTTT CAACTGTCTA AGCAAAGGCG TGAAATATC	4860
TGTACTCAAG CCTTCTAAAA GCTTGCTGGC TACTTCTACT TGATCGATAA TCTTTCTGA	4920
TTTCCCCTGT TCCAAGGCTT CTACCAGAGA AGTCACCGTT TCTTTGAGG AAGTTAAAAA	4980
ATTTTGATTG ATATTTGCT TGATTTGCTG GACCATGTGA CTCGATACAG CCACCTCCTT	5040
GGTCCATCCC ACTAAGAAAT CACATTCTAA AGTTGGTTTC ACTTGTGAAA TTGAAAAGCC	5100
CCAATCACGC TCCAGAACTG TCGCCAAGTT TTCTTCTTCT AACCAAGCAG CCACCTCTG	5160
GCGATCAAAT GACTGGTAGA GAACCAAATC CTCTGCCACA ATACAGGCAA GGTCGCCCAT	5220
GGAACCATTG TCTCCTCGCT TAAGCAAGAC AGCGCTAGTC AGCTGAACA AGAGCTCCTG	5280
ATCAACAGAA ACATCATACA GAGCCAGTAA AGCCTTGACA ACCAAGACAA CGACGCTGCC	5340
ACTAGAACCT AGACCAAAC TTTTCCCTTC TCGTTCCATT TTGCCACAGA TTTCTAGAGA	5400
AAAAGGTCTT AAATTCTGAC CACGAACAGC GAGGAAGTCT CCCATCAAAG CAATCGTTTC	5460
TTGAATCAAG CTATAGTCAG GATTAGGCCT TAAGTCCACT GCGAAATCAA ACATATCTGA	5520
ATAGATAACGG TAGCTGTCAG AAAAGCAAT CTCAGCCCTC ATATAGATGG GAATATCCTT	5580
TATCAAAGCT AACTGCCCTG GCTCTAAAAT AGCATATTCA CCTGCCAAT AGAGTTTCC	5640
GCAAGTTTA ACAGCAATCA TCTTGACTCA AATCCTTTGT TTTTGACACA ATCAAGCGAT	5700
AACGATGACC GAAAATTCT GATAAATGCT CCAAGTCTTT CTCCTGACAG AAGACCTTAA	5760
CATTGGGACC AGCATCCATG GTAAAGTAGC AGGCCTCTCC TTTCTCACGA AGCTGGCGAA	5820
CAAAGGCCAT AGCCTCATAA GAGGCATCCG TCAGATAAGA AAAGGCTGGA CTAGCAGTCT	5880
TTGTCGTAGC ATGCATAGCC AGGGCATTTC TCTCCGTTAA TTCTCCAATC TTGGCAAAAT	5940
CATTTCCCTT GAGATAAAC AGCATATCCT GATAGTCCTT CTCAGACTGA CGAACCCAGT	6000
CGTCGAAAGT CGTCGAGGTT TCCACACAAA GTTTCATCCC GTCACGGCTA GAGATTGGTT	6060
TTTTCTTGTC CTCTAGCACC AACATAATCA TAGCTAGTT CAAGTCTGTC TCTACAGGGT	6120
AAATTTCTCC ACTATCCCTA TCCCAGGCTC CTAGTGGTCC ATAAAAACTC CGAGAAGAAG	6180
AACCTGAGGC AAATTTGGCT TCCTGTGCCA ACTGACTTCT ATCCAATCCA AGCTGAAAT	6240
AAGCATTACA AGCCTTGACC AGGGCGGACA AACCACTAGA ACTTGAGGAC AGACCCGCTG	6300
CCGTAGGCAT ATTGTTTGA GTATCGATAC GGACAAAGCC CTCACCGAGCT GGACGATAAC	6360
GGTCAATAAT CTTACTCATC TTGGCATGCT CGACCTCATT TTGTAGCTGA CCATTGATGT	6420
AAAATTCGTC AGCTGTTACA TTGGCTGGTA AAGGCGACAA GGTCGTCTCT GTATACATAT	6480
TTTCCAAAGT TAGAGAAATA CTGCTAGTAG CAGGCACCAT CTCTTTCTT TTTTCTTTC	6540

982	
CCCAATATTT GATAATAGCA ATATTTGCGT AGGAACGTAC TGTTACAGGC TCTCTATCCA	6600
TGTCTGAACA GCTCCTTTCT CTTCTAATCT TTCTGCTAGT TCTTGTCGCGT GTGTCAAATT	6660
GGTTACCAAG GCTATGATAC AACCTCCTAG CCCACCACCG CTCATCTTGG CACCCAGAGC	6720
ACCATGGCTA AGAGTCGTTT CAACCAAAAA GTCTGCCTCA GGGCTACTGA CTCCAATTTC	6780
TTTTAAATGT AAATGCGCTT GACTGAGGAT TTGTCCCAGT CCTTCAGCAT CTTTTGTGA	6840
AATCGCAACT TCTGCTGCT GGGTTAACCTC TCCCAAGGCA TGCAAAAACG GTAGGGCATHC	6900
CTTGCCCTTA TTTTGAACCA CTTGGATGGC TTCACGAGTA TGACCATAAA CACCCGTATC	6960
GGCAATCACC AAATAGGCCG ATAAATCCAT CTCAAGTTCT GTAAATCCTA CGTTCTTGAT	7020
AAAGCGAATA GGTTGGTCAC TAAGACAGGT CTTAGCATCC AAACCACTAG GATTCATATG	7080
GGCAATCATT TCAGCTCGAT TGACCAAGAT TTCTAGTACA TCATGAGGCA GATCAGCCTG	7140
ATAGTAGTCA AATACTGCAC GAATGGCCGC TATGCTGATA GCCGCTGACG AACCCATCCC	7200
CCGTTTCTCA GGGATAGCCG AGTCAATCTC ACAACGAATG CAGGCTTCTG TGATATTCAA	7260
ATACTCCAGT GAGGCATAAA CCGCCATGGA CAAGGTATCC TCCTCATAAA GGCGCCAAGG	7320
ACTCTCTGCA GGAACCTACCT TACAGGTACAC CTCCACCTCC AAAAGAGGCA GGGAAATGGC	7380
AGGATAACCG TAAACGACCG CATGTTCCCC TATTAAAATT ATCTTACTAT GTGCCTGACC	7440
GACACCAACT TTTTTGTCA TTTTTCCCTT TTACTAGACG AAAAAACGTC TTATTTTCA	7500
TACAAGTATT AATTCTTCC TATCTATTTT ATTATATTTT CACAAAAAAA GCGATTGTTT	7560
CCATTCAACAA TCGCTTCTTT CATTATTGAA CCCATTCGCC ATTATAGTTG ACAGAATAGC	7620
CATCTACGGT CGTATTCACT GCCAAGGCAC CTGAGCGCTA TAAGCGTAGT ACCATCTGCC	7680
ATTGACCTGG AACCAACCTG TCGTCATAGA ACGACGAAAG AAACCTCCATA CCATTAAGTA	7740
AAGAGGAAAG TCGTGAGGGA GCATGCGCCA TTGACAACCT GTTTTAGTGA CGTACAAAGT	7800
CTCATTAACA AGTACTCGTT TCGGCCATTT ATAGGTGCGG TGTTTGGAGA AATAGGGTTC	7860
AATCTTCGCC CATTCTTGAT CGTTAAATC AGTATCATAT GCTTTGCCGA TCATAACTCT	7920
AGCTTAACAT TTTTTGTGA ATACAGGTTC TAAATAATCG ACCACGAAAA TTTCTTAAGT	7980
GGAAAACGCC TTATGAAGTA TGCTACGGGA AAGTTATGCA CTTAATTGA CAATTCAAGA	8040
TGTAAAATA TATACTATAG TAGATTGAAA CTAGAATAGT ACACCTCTAC TTCTAAAATA	8100
TTGTTAGAAA TCGATTGAC TGTCTGATC GATTTATCCT GTTATTATCT CATTCTACTA	8160
TAATATTTGA TAAGTTATCC TAAAAGTATT ATTATGTTGT TGTGTTATAG ATTGATTGAA	8220
TCTAACTAAA GGATCCTATT CAATTACTAG AACTATCACA TACTCAAGGT CAGCTCACAG	8280
ATGAGCAACT ATTTGGTTA CAATGTCTAC TAAATTAAAG TCAAACAAAT AATTTAGTCA	8340

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AAATTAAAAA AATAGAGGAA CATAAATATG ATTACAAAAC AGAATGTAAT AGTGTCTAC	8400
AATTTTACT AGATAAAACT GTAAATTCTG AAGGAAGGAT CACTTCTCA ACAGAATTG	8460
GAAATTCGT AAGTAATTAA TCATTCCAAC ACGGAAATGC TGGACTACTG TTTCCTCTAA	8520
ATAAATTGTA CCCCCCAGAA CTGGATTCTA AAATACTCTC TATCATCAAG AAGGCAGTGA	8580
CAATTAGAAC GACACACACA TATGAATATC AATACTCACT GCTATTTGGT GATGCAGGCT	8640
ATCTATGGTT ACTCCTACAT TTATTTCTA TCAGTAAAAA TCAACTATAT CTACAATTAG	8700
CAAACGTCAC CGCTAAAAAA TTAATAGAGA ATTATGATAC TCTAGAGGAA ATAGACTTTG	8760
CATTGGGAAA ATCTGGTGTCT ATTATATCAT TAATAAAATA CTATCAATT ACCAATGACA	8820
ATACTCTTAA AATTTTCATC CACAATAGTA TAGGGGAAAT TTATCATTAT TTCCTACAAA	8880
GAGATACAGC CAAAGAAAGC ATTTAGACT ATAGCTTGCG TCATGGATAT TGTGGAATTG	8940
CATATGCTTT ATTTGCCTAT TCTAAAGTCT TAGAACCTTC TATGTTTAT AATGATCTCC	9000
ATACATTCCA TACTGAATTA AAAAATTAT TAGAAAAAGT TACTCTAAT ACTGAAAATT	9060
TAGGAAATT ACAACTTCT TGTTGCAAAG GAATTCCGG AATAATCTTA TATCTTGTA	9120
TGTACGATTG TGACGGAAAC AAAGATATTA TTAGTAAATA TCAAGAATTG GTTTTAACC	9180
ATCATCTAAA AATGATGACA GGATATTGCC ACGGAAATAAC TAGCTTACTA CAAACCACTG	9240
TCTACAATCA AAACAAATTA CTGATGAAAA AAATCCAACA GGTAATTAA GCATGTTCTG	9300
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ACTTCGGAAT AGGAAGCATG GGGTATATTG GTGTCTATTA AATAATAAT TCCCATTGCA	9420
TGTGCAGACA TAAGGAGAAA AGTATGAAAT TATTTGGAC AAACAACATA TATAGACAGT	9480
TGCTGCTAAA CAGCTGTTT TCATCATTGCG GCGACAGTAT TTTCTACCTC GCCATTATCA	9540
ATTATGTGGC TCAGTACAAT TTGCTCCGC TAGCGATTTC ACTGATTTC ACTTCAGAGA	9600
TGGTCCCCCT ACTATGCCA CTCTTCTCG GGATTCTAGG AGATTTCAA GAAAATAGAG	9660
TCAAACACGC ACTCTGGATT GCCAAAATCA AAATCCTGCT CTACGCTATT TTGACAGTAT	9720
TTCTCGTCTT GTCGCCCTTT TCATTAGTTT CAGTCATTAT GATTGTCATC ATCAACCTCA	9780
TCTCTGACAC CTTGAGCTAC CTGCTGCCT ACATGATGAA CGCCCTCTAC ATCAGTGTAA	9840
TTAAGGACGA CCTGCATGAT GCCATGGGGT TCAGGCAGTC TCTGATGAGG GTTGTCCGTA	9900
TTGTCGCCAA TCTGGCTGGC GCATTCCTTA TCAATGTTAT AAGTATTCAA ACTATTTCCC	9960
TTATCAACAC TCTGACTTTT GTCATTGCCT TTTTGGGCCT GTATGTTATT CGACATACCT	10020
TGTATGAGGT TGAAAAAAGA ATTGAAATGT CACATACAGC ACTGAGTTT AAGAAATATT	10080

984

TTCAACATCT	TAAACAGTCG	CTGGCTGTGC	TCCTGAGGTT	AAAAGATACC	GTCATACTAC	10140
TGTTTCTGAC	GACCAGTATG	ATTGCCATCT	TGGATGTGTC	CCCTCGGCTG	ATTGCCCTCC	10200
GCTTCATCCA	ACAGACACTA	GCACAACCTGA	GCATTGGGCA	ACTCCTCGCC	CTGCTCTCCA	10260
TCATCATGTC	TTGTGGAGCT	ATCCTTGGCA	ATATGACCAG	CAGTAATCTA	TTTAAAAATA	10320
TCCGTTTCAC	GCACCTCTTG	GTGTTCTGTG	AGATTTCCCT	ATTGACTCTA	ATAACTAGTA	10380
TCCTTTGTCA	AGCCTATATC	GTAATTTCA	TGACCAGTTT	CATCACTTCT	ACGATTATCG	10440
GCATTCTCAG	CCCTCGCCTA	CAAGCAGCTG	TCTTGCCCA	TATCCCCAGT	GACAAGATGG	10500
GGACGGTGGG	CTCTGCTCTG	AGCACAGTGG	ACATTCTCGC	CCCGTCCCTG	CTCTCCCTAT	10560
TAGCCCTATC	CATAGCATCG	GGCGTTTCGG	TGCAGTTAGC	ATTGATATT	TTGTATCTTA	10620
TTTTAATTGC	TCTTATCTTT	TGTCAATGGT	TAGTCAAGTT	CAACACTCAT	AACTAACGAA	10680
AAAGCATGTG	TAGATTCAC	ATGCTTTAA	TCTCCCCAAT	CGTCAGGTCA	AGTACAACAA	10740
AGTCACTTCT	TTGATTAAGC	GAGTGTCTA	ATATAATTAT	AAGCGCCCTG	TCATTACCGA	10800
ACCCATTCCGC	CATTATAGTT	GACAGAATAG	CCATCTACGG	TCGTATTAC	TGCCAAAGCA	10860
CCTGAGCTAT	AAGCATAGTA	CCAGTTGCCA	TTGACCTGGA	ACCAACCTGT	CTTCATGTCT	10920
CCATTACCTG	CATTTAGGTA	GTACCAAGTT	GAACCATCTT	GATACCAACC	AGTTGCCATA	10980
GCTCCTGATG	AACGGAGATA	GTACCATTTG	TTCCCAAGGT	TTTGCCAACC	TGTTTCATA	11040
TCGCCATTTG	GGTGGCTAA	ATAATACCAA	GTGGTACCTT	CCTGATACCA	GCCAGTGGCC	11100
ATTGCTCCTG	AGGAACGGAG	GTAGTACCAAC	TTATTACCTA	GATATTGCCA	ACCTGTTGC	11160
ATAATACCAAG	TTGTTGGATC	TAGGTAGTAC	CAAGTCGAAT	CATCGTTAT	CCACCCCGCA	11220
CGTCTTCAC	CACCAAGGTA	GTCTTCTCCA	TTAATTTCCG	TCTTAGCTAG	ATAATACCAAG	11280
TTAGACTGAT	CATAAAGCCA	ACCTGTCTCT	AAAGAATGAT	TTTGATTAAA	GTAATAGTTC	11340
GTATAATAAC	GCTCTCTTC	TTTATCTTCT	GAATCTTCAC	GTTTTCCCC	GTACTTTCTT	11400
CCAACACTGT	CTTTAGTTTT	AATCTCTAAT	GTTCCTAAC	CAACAAACTC	TTGTAGCACT	11460
CCATTTTTAT	CGAAGTAGTA	CCACTCTGAC	TTTGGAAAAC	CTTCTAATCT	GATACCATT	11520
GGGTAAGGAC	CAATTGTACT	ACCTTTAGAT	GGAAACGGGA	TATATTGCCA	GCCGACAACC	11580
ATCTCTCCAG	ATAGAGAAC	AAAATAATAG	TACTTACCAT	CAATCACTCG	CCAGTAGGTT	11640
TCTTGAGGT	CCCCCTTTTT	GTAGTAGGTT	CTTCCGTTTT	CTTGGACAAA	CTGCCATCCT	11700
TCAGAACAT	CTGCAAATAC	TGTACTGGTC	CCTAGCAAAC	CAAAGAAAAA	TACTGTCAGT	11760
CCAACATTGCA	TAGTTTTTT	CAAATTTTC	ATCTATATAC	CCTCCAATAT	TAAATCCACT	11820
CACCAAGATGA	GGCGAAATTA	TAAACTTTAC	CATCGATAGT	TTGGCTACCT	GTAACCATTG	11880

985

CTCCAGG

11887

(2) INFORMATION FOR SEQ ID NO: 147:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 11340 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 147:

CCGGTATGTT CTGGAATACT ACCAATCTAA GCTGGCTGTG CCCTACAGTT TTACAACCT	60
GTACGAATAC CTTAAGGAAT ATGACCGATT TTCAGCTGG GTTTGGAGT CTGGTATTTC	120
AAACGCTGAT AAAATATCCG ATATTCTTT ATCAGTTTG GAAAATATGT CTAAGAAAGA	180
CATGGAATCC TTTATCCTT ATCTACGTGA ACGTCCCTTG CTGAATGCTA ATACAACAAA	240
ACAAGGTGTT TCACAGACAA CTATCAATCG AACCTTATCA GCACTTTCTA GTCTTACAA	300
GTATCTAACG GAGGAGGTTG AAAACGATCA GGGGGAACCT TATTCTATC GTAATGTAAT	360
GAAAAAAAGTT TCCACCAAGA AAAAGAAAGA AACCCTTGCT GCCAGAGCTG AAAATATCAA	420
GCAAAACACTC TTTCTAGGTG ATGAAACAGA AGGTTTCTA ACTTATATCG ATCAAGAGCA	480
CCCACAAACAG CTTTCAAATC GAGCTCTCTC ATCATTCAC AAAAATAAAG AACGAGATT	540
AGCCATTATT GCCCTCTCT TGCGCATCTGG TGTCGCTTA TCTGAAGCTG TTAATCTAGA	600
TCTAAGAGAT CTCAATCTAA AAATGATGGT TATTGATGTT ACTCGAAAAG GTGCAAACG	660
TGACTCAGTC AATGTCGCTG CTTTGTCTAA ACCTTATTTA GAGAATTATC TGGCATTG	720
GAATCAACGC TATAAACCGG AAAAACAGA TACAGCCCTT TTTTTAACCTC TCTACAGAGG	780
TGTTCTTAAT CGTATCGATG CTTCTAGCGT TGAGAAAATG GTTGCTAAAT ACTCAGAGGA	840
TTTTAAAGTG CGTGTAAACAC CCCATAAAACT GCGCCATACA CTAGCAACTA GGCTCTATGA	900
TGCGACTAAA TCACAAGTT TAGTCAGTCA CCAACTAGGA CATGCTAGCA CACAAGTCAC	960
TGACCTCTAT ACCCATATTG TTAGTGTGAA ACAAAAGAAT GCTCTGGATA GTTTATGATT	1020
TTACGTATTT TAAATTATGT AAATAATAT CAAAAAAAGA AGTTGGCCAA CTTCTTTTG	1080
ATTTATCCAA CTACCGCTTC AGCGATTTCT TCACGGCTAA TACCAGCGAA GTAGCGTGTG	1140
ATATCAATGG TTTTAGCGC CTTAAGAACAA TCTTCGCGTT CGTATTCAC CCCACGAAGG	1200
ACATCTTCTA CTGCAGCAAC GTCTTCATA CCAAAGAAGT CACCATAAAT CTTGATGTCT	1260
TGGATTTTG ATTCAAGAAC GTTAGCAAAG ACTTCAACCT TACCACTAGT GAATTTGATT	1320

986

CCACGACGGA CGTTAAATTC AGGTGATTAA CCATAGTTCC AGTCCCAGT TCCAAACTTA	1380
GTATCCTTGA TGCGATTGAT TTGGCCAAT TCTTCTTCTG AAAAGACGTA TTCAGTCATC	1440
TCTGGGTACT CTTTTTCAT GTATTCCAAG AGTAAATCAC GGAATTTC GACTGTGATT	1500
TTTTTGGTA ATTCATTGAT AATATTGGTT ACACGGGCAC GGACGGATT CACACCTTT	1560
GATTCAAATT TATCTTTGA AACCTTAAGG GCATTTGCGA GGACTGACAA ATCAACGTCA	1620
AAGAGCAAGC AACCGTGGTG CATGATACGG CGCTTGATAT AGGCTGGGC ATTGCCACAG	1680
AACTTCTTAC CATCAATCTC AAGGTCATTA CGACCTGTGA ACTCAGCTT AACCCCAAGT	1740
TGAGCCAGGG TATTGATAAC CGGAGTTGAG AAGCTCTTGA AGTCAAATGC CTTATTTCA	1800
TCTTCTTGG AGATGATCGT GTAGTTGAGG TTATTTAAAT CGTGGTAAAC AGCTCCACCA	1860
CCACTAATAC GGCAGACTAC CTCAATACCA TTTTGCAGAA CATAATCACG GTTGATTCT	1920
TCGATAGTGT TCTGGTGACG ACCAACAAATG ATAGATGGCT TGTAAATCCA AAGTAGGAAG	1980
ATTTGATCCT CATCCAAAAG GTGTTAAAG GCGTATTCTT CCAAGGCAAT ATTAAAAGCA	2040
GTGTCATTG AATGATTGAT AATGATTTTC ATGATATCCC TTTACTTTAT ATGATAGAAA	2100
CTGGAAATAA CCTTCCAGTC TAATCTATCT TCGTTTTATT TTTTCTTAGG TGAATGGATG	2160
GCCATTCCCTA GAACATCTGC AAACGCTTCG TACATCACTT CAGAGTAAGT TGGGTGCCCG	2220
TGGATGGTCT TCAGCATTTC CTCAACAGTG ATTTCCATT CGATGATGCT TGATGCTTCG	2280
TTTATTAATT CTGCGGCTGC AGGACCAATA ATGTGTACAC CAAGGATTTC TCCGTATTTC	2340
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GCAGCAAAGT TAAACTTACG GATGGCAACA TCGTATTCT CACGGGCTTG TTCTTCTGTC	2460
AAACCTACTG CTGCTACTTC AGGGAGAGTG TAGATGGCTG CAGGAGTCAG ATTCAATTG	2520
GCAACTGCAT GATTCCTTT AAGGGCATTT TCAGCGAAA CTTCACCCAT GCGGAAAGCT	2580
GCGTGAGCCA ACATCTTAGT ACCGTTGATG TCACCTGGTG CATAAAATGCC TGGAACTGAA	2640
GTTTCCATGT ATTGTTGAC CTTGATACAA CCACGATCCA ATTCAAACCTC AACCTCTCCA	2700
ATACCTTCAA GGTCTGGCAT ACGACCAATT GAAAGAAGAG CTTTGCTTGC GATGATATCG	2760
TCTTTCCATT CAACCTTGAT ACGAAGTTGA CCATTTCTT CAATGATTTT TTGCAGTTA	2820
GTACCAGTCA AGATGGTCAT TCCTTACGC TCAAGAATCA AGCGAAGGTT CTTAGAAACT	2880
TCCACATCCA TAGCTGGAAC TATACTGGTCC ATCATTTCGA TAACAGTCAC TTTGAACCA	2940
AATGTCATGA AGGCCTGACC GAGTCGATA CCGACAACTC CACCACCGAT GATAACAAGG	3000
CTTTCTGGCA CTTCGTTCAT TTCAAGAATG TCATCACTAG TCATGACAAG TGGAGATTCC	3060
ATACCAGGGA CGTTGATCTT GTTGACTTTT GAACCACCAAG CAAGAATGAT TTTCTTGGTT	3120

987

TCAAGCAATT CAGAACCAT TACCAAGACG TTCTTGTCTT TAGTGATTGT ACCAATTCC	3180
TTATGAACAG TAACTCCGTA GCTACGAAGA AGTCCTGCAA CACCACCAAC AAGAGTATTA	3240
ACAACCTTAG ATTTAGTTTC TAAAAGTTT TCCATATCAA CAGTGAAGTT AGGATTTCA	3300
ATCACGATAAC CACGATTGC AGCATGACCG ATATTTCAA TAATTTCAGC GTTATGAAGG	3360
TAGGTCTTGG TTGGAATACA TCCACGGTTT AAGCAGGTT CACCAAGTTC AGATTTCTCA	3420
ACAAGGGCAA CCTTACGCC GAATTGGCA GCTTTAATGG CTGCAACATA ACCAGCAGGA	3480
CCTCCACCAA TCACAACGAT ATCAAAAGCA TCATCGCTCT TACCATCATC GTTGAGGTA	3540
CTTGCTACAG GTACAGGGCT AGCTTCTGGC GATGCTGCTC CAGCTGTTGG GATGTTTCC	3600
CTTTCTTCAC CAAGGTAACC GATAACTTCC GTTACAGGGA CAGTTTCAAC ATCTCCTTG	3660
AGAATGGCAA TCAAGTACCC ATCTTCTTCG GCTTCCAATT CCATGCTGAC TTTATCAGTC	3720
ATGATTTCCA AAAGGATTTT TCCTTCTTT ACAAAATTCTC CGACTTTTTT ATTCCATGG	3780
ACGATTTGTC CTTCTGTCAT ATCCACGCCG GCTTTGGCA TAATTACTTC TAAGGCCATG	3840
TCTTCCTTCC TTTATCTATA TCTTAAAAAT GAATACTCTT GCTCTTAAAT TAACATTGAG	3900
ATTGGCGTTT CAATCAACTC TTTCAAGTCC TTCATAAAACT TAGCACCAGC CATACCAC	3960
ACGACACGGT GGTCAATGGT TAATCCTAAA CTCATGATTG GGCGAATCAC AATTCACCA	4020
TTGACGACAA CTGGCTTCTC GATTGTCGAA CTGACACCAA GGATAGCTGA GTTGGTTGG	4080
TTAATAATCG GACCAAGGA CTGAACACCA AACATTCCC AACATTGAT TGTGAATGTT	4140
GAATTTGTA ACTCACTTGG AGCCAATTAA CCATCCAAGG TACGGCCAAT AACATCCTTA	4200
AAGGCTACAA CCAGTTCTGA AAGACTCATC TTCTCAGCAT TGTAACAAAC AGGTGTCATC	4260
AATCCATTAT CCATCCCAAC TGCCATGGCA AGATTGACAT AGTTGTGAGT GATAATAGTC	4320
TTGCCATCTT CTGTCAATGA AGCGTTGATG TATGGGTGTT TCATAAGAGT CTTAACAACT	4380
GCAAGCGAAA GAAGGTCTGT TACAGTAGTC TTCTTCCCAG TTGCTTCCAT GATTGGCTCA	4440
AGAACCTTCT TACGAAGAGC CAACATTCA GTCATATCAA CTTCATAGTT GAGGGTGAAG	4500
GTTGGCGCAG TCAAGTAAGA TTCAACCAG CGTTGGCAA TAACCTTACG CATTGGTGTC	4560
ATTGGAATAC GCTCGATTTT ACCATATGGT GTTACGTTAT CAGGGACTTC TTCCACTTT	4620
TCAATCTGAG CAGGAGATTT GATGCTATCG TTTCGATAT TTTCAGGAAG CAGGGCCAAA	4680
ACATCCTTCT TCATGATTTT ACCACGATGA CCGGTTCCCTT GGATTTCTG CCAAGCAATG	4740
TTATGTTCGA GGGCAATTG TTTTGCAAGT GGCAGAAATGC GAACCACGTT TGTGTCTTA	4800
TAAGTTCCA CGTCTTCTT GTGGACACGA CCGTTGCAC CTGAGCCAGA AACGTCGTAG	4860

988						
AGGTTTATCC	CTAAATCATC	CGCTAACCTT	CTAGCTGCAG	GAGTCGCTCT	TAGCTTGTCA	4920
TCAGCCATGA	CCTCTCCAAT	TCTATTATG	ATACAAAGGG	CGTAAAAGC	GACTGAAAAA	4980
TAGGAAATCG	ACGATGGCTT	CGATGAAGCC	AAGGAGATTT	ATCTTTTTC	CGATCTTTA	5040
GCCC GTGCTC	TAATCTAAGA	TATTAATGAC	GAAGAGCTCT	GCACCTAAA	GATACAAAGT	5100
TTCTCGTCAG	CTTTATTTA	TTTACATAAC	TTATCTTATG	TAACCCTATT	CTTTGTTATA	5160
AGTTTTTCGG	ATTGCATCTT	TGATACTTTC	AACTGTTGGA	ATCATTGCAT	TTTCTAGGTT	5220
TTGTGCATAA	GGCATCGGCA	CATCTCTCC	TGCACAACGG	CGAATTGGTG	CATCTAGATA	5280
GTCAAATGCT	TCTGATCTG	AAATAATAGC	TGAAATTTC	CCGATATAGC	CACTTGT	5340
GTGGGCATCG	TTGACCAGAA	CAACCTTACC	AGTCTTCTTC	ACTGAGTTA	TGATGATATC	5400
CTTATCAAGC	GGAACAAAGGG	TACGTGGT	ACAATTTC	ACTGAAATTC	CTTCTCTGC	5460
TAATTCTCA	GCAGCTTGAA	CCACACGGCG	AAGCATT	CCATAAGTAA	CAACTGTTAC	5520
ATCCGTTCC	TGGCGTTGA	TTTCACCAAC	CCCAAGTGG	ATTGTGTAGT	CTGGATCAAC	5580
TGGCACTTCC	CCTTTTG	TAAATTCTGA	CTTGTACTCA	AGTATAATAA	CTGGGTTGTT	5640
ATCACGGATA	GAAGACTTAA	GCAGGCC	CATGTCCGCA	GGTGT	GTGCCACAAAC	5700
CTTAAGTCCT	GGAATGTGAG	TAAACCAAGA	CTCTAGAGAT	TGTGAGT	GCGCAGA	5760
GCCAACCTCG	TTACCAGCTG	CACAACGAAC	AGTCATTGGA	ACCTGAC	TTACCAACAA	5820
CATGTAACGT	GTTTAGCAG	CTTGGTTGAC	GATATTGTCC	ATGGCAATAA	CAGAGAAGTC	5880
CATGAAGGTC	ATATCGACGA	TTGGACGAAG	TCCTGTCATG	GCTGCTC	CTGCTGCTCC	5940
AGAGATGGCA	GCTTCAGAAA	TCGGACAGTC	ACGGACACGT	TCTGGAC	AAATTCTCAAG	6000
CATTCCAACA	GAAGTACCGA	AGTCTCCTCC	GAAGACACCG	ACGTCTCTC	CCATCAAGAA	6060
CACATTTCA	TCGCGACGCA	TTTCCTCAGA	CATAGCAAGG	ATAATGGTGT	CACGGAAGGA	6120
CATTGTTTT	GTTCCATT	TATCTCTT	TCCTTAGTCT	GCGTAAATAT	CTTCAAAGGC	6180
TGATTCAAGC	GGTGGGAATG	GGCTTCC	TGCAAATT	ACAGAAGCTT	CTACTGCTC	6240
CTTTACTTGC	GCTTGGATT	CTTCCAATT	TTCGGCAC	GCAATGTT	TTTCAATAAG	6300
GTAATTGCGG	AGGTTTCGA	TTGGATCTT	TTGTTCCAC	AATTCCACTT	CTTCACCGT	6360
ACGATATT	CCAGGGTCAG	ATGATGAGTG	ACCGAGCCAG	CGATAAGTTA	CACTTCAAT	6420
CAAGACTGGA	CCATTGCCAC	TGCGAACATG	GTCCACAGCT	TTCTGAAATC	CTTCATAGAC	6480
ATCGATGACA	TTGTTACCGT	CTTCGATGAA	CATTCCAGGA	ATTCCATAAG	CGGCGCTACG	6540
TTGATGGATA	TGTTCTATAT	TGGTCATTT	CTTGATATCC	GCAGAGATA	CGTAACCGTT	6600
GTAAATGCAA	TAGAAAATGA	CTGGCAGGTT	CCAGATAGAA	GCCATGTTCA	CTGCTTCGTG	6660

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GAAAACACCT	TCATTGGTCG	CACCATCTCC	AAAGAAGCAG	ACAACGATTT	TACCGGTATT	6720
TTGCATTTGC	TGACTGAGGG	CTGCACCGAC	AGCGATCCCC	ATACCACCAC	CTACGATACC	6780
ATTGGCACCA	AGGTTCCCAG	CATCAAGGTC	AGCGATATGC	ATAGATCCAC	CTTCCCTTT	6840
ACAGGTTCCA	GTGTATTTAC	CAAGGATTTC	AGCCATCATT	CCGTTGAGGT	CAATCCCTTT	6900
AGCAATAGCT	TGCCCGTGTC	CACGGTGGTT	TGAGGTAATC	AGATCATCTG	GATTGAGAGC	6960
TAACATAGCC	CCCACGTTAG	CTGCCTCTTC	ACCAACAGAA	AACTGCGTCA	TTCTGGCAC	7020
TTTCCCTTTC	TTTACTAATT	GTGCAATTTC	AAAGTCCATG	CGACGGATTT	CTTCCATCTT	7080
ACGGAACATT	TCTAGCAAAA	GATTTTATC	AAAGTTGAC	ATCTTCTTGC	CTTCTTAAC	7140
TTCTTCTTAC	CTTACTATTT	TACCGCTTTT	GGCAAATACT	GTCAAAGTTT	TTCTAAAAGA	7200
AATTTCACAA	AATAAAAAG	AAACCCCGT	AAAAACAAGG	GATTTTCTTG	TCAAGAATAT	7260
TTTTTCACAA	ACTTTTTAGC	ATTTGGATTT	TGCTAAAGAT	TCAAATCTCT	TCATAATCAC	7320
AGTTAAACCG	CAACGGTAGA	GCGCCCCGCT	CACAATCAA	CTAATAATCA	AGCCGATCCA	7380
GTAAGAATAA	GCTCCAAAAT	CTGTTAGGGA	ATCAAATAGC	GTAnCACAGG	GATTGCTACG	7440
CCCCAATAAC	CAAGCAAACC	AAGGTAAAAA	GGAATAACTG	TATCCTTATA	CCCCCGCAA	7500
ATTCCCTGAA	GCGGCGCCGC	AAAGGTATCT	GCTAACTGGA	AGAAAAGACT	ATAAGTTAAA	7560
AAACGCACTG	TCAAATCGAT	AAATTTGGG	TCGTTACCAT	AAAGACTGGC	CACATTCCC	7620
CTAAAAATGT	AAAGGAAGGT	TAAGGTGAAG	GCGCAAAAA	TGAGGGCAGT	CCATCTTCCT	7680
AGACCAATAT	AGGTTTCGC	ATCATCAAAT	CGCTTGGCTC	CCACTTCATA	GGAAACGACA	7740
ATAGCCATAG	CCGATGAGAT	ACTCATAGGA	AAGGCGTACA	TAAGACTTGA	AAAGTTCTATA	7800
GCTGACTGGT	GACTAGCTAT	AATCAAGGGC	AAAAACTTAG	CCATAATCAA	GCCAACCACT	7860
GAAAAGATAG	CCACTTCCGC	GAAGACAGTT	CCCCAATAG	GCAGACCTAA	ACGAACTCCT	7920
TCCTTAATT	TATCCATATT	AAGTGGATT	CGTTTCTCAA	GGTGTAAGGC	TTTGAGCTTC	7980
TCCTGTTAA	ATAAAACCAG	AACAGAAATC	CCAAGCAAGA	CCCAGTAGGC	CAAGGATGTT	8040
CCTAAACCAG	CACCAGCCCC	TCCCAGTTCT	GGAACACCAA	AGGCACCGTA	AATCAAGAGA	8100
TAGTTAAATC	CGCTATTGAG	AGGGAGTAAC	AAAAGCATGA	GGTACATGGA	CAGTTGGTC	8160
AAGCCCAGCG	AATCCAGCAA	GGAACGAATG	ACGCTAAAGA	GCAACAAGGG	GATAATCCCG	8220
ATAGATAAAA	ACCAAAGATA	GCGAACCGCT	ACTGCCGCTA	CTGCTGCTTC	TAACCCAATA	8280
TGATTCAAGA	TTATTGGTGC	CAAGAAAAGT	ACCATCCCCA	GCAAGACCAC	AGATAGGCC	8340
AAGGCCAAAT	AAATAAATTG	GTAAAATCA	GACGCAACTT	CTTCCTTTT	GCCTCGACCA	8400

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AGATGGTGAC CAATGATAGG CACCAAGGCT GACACAATCC CTGTTAGAAA TGTAAAGAAA	8460
GGATTCCAGA TACTGGTTGC CATAGATACA CCAGCCAAGT CCATAGTGT GTATTGACCT	8520
GTCATTGCAG TATCAACAAA AGAGGCAGAA TAATTGGCAA ATTGGTAGAT CAGGATTGGG	8580
AAGAAAATTT TTAAAAATAA TACTAACTTC TCTCGTAAAC ACTTTGTCTT ATACATACTT	8640
CTCTTTCTAT TCTGATTAT CTAAACCAA GAGTTTCAGA CCATAGTTT TCAAACCTAG	8700
CGGAGGTTA TTAGATTTC AAGTAGTATG CCAACACGCA CATGTACGAC AATAATAGCT	8760
TCTAACTAAA CCTCCGTTAT CATATTGAAC CGCATGGTCA GCTTTTCTT TAGTTTCATA	8820
TTGAATTTC GAACGATTAG CTGCGGGACA GTAAATTCCA CTATTAGATT TCGCTTGTCT	8880
CTCCCTACGT TTTCGAAAAT AATTCACTATT CTAACCTCCTA TCAAGCTTGA TAGACGATTT	8940
GTCCCTTACA GATGGTATAT TTAACCTGCC CTTTTAAGGT TTCACCGATG AATGGTGAAT	9000
TAGCTGCTT GGAAGCAAAA TGGGAGTCCA CAAAGCGGTC AGCCTGGCA TCAAAAATAG	9060
TGATATCTGC TGGACCATTG TCAGCCAAGT AACCTGCTTC AAAGTTGTAA AGCTTGGCTG	9120
GGTTGTATGT CATTTCATCA AGTAATTCCA TCAAGCTCAA CTCACCAGCT TCTACTAAAT	9180
AGGTCAAGCT GAGAGACAGG GATTTTCTA AGCCAGTCAT ACCAGATGGC GCTTTGGTAA	9240
TATCCTCAAC ATTTTTTCA TCTACATGAT GAGGCGCGTG GTCAGTCGCA ATAACCTGTGA	9300
TGACACCTGA TTTGAGACCT TCGATAACGG CACGACGGTC TGATTCCAAA CGAAGCGGTG	9360
GATTCTCATCTT AGCATTGCTA CCTTGTGTTA AAAGAAGTGC TTCTGTCTTA GAGAAATGCT	9420
GTGGCGCTAC TTCTGCTGTG ACTTCTGCAC CTAACCCCTG AGCAAACCTCC ACTACTTTAA	9480
CACTTCTTC CTTAGACAAA TGCTGGATGT GAACATGGGC TTTAGTTGCA TAGGCAATCA	9540
TGACATCACCG CGCCATCATA GCGTACTCAG CCACCCCCAGT AGCACCGCAG ATATGAAAT	9600
GTTCTCTAGC AATATTTCA TTAAAGCCAA GAACACCGTT CAAACCTGGA TCTTCCTCAT	9660
GAAGGCTGAT AAAGGTATTG AGTTTTTGG CTTCCCTCCAT GGCTTCCTTG ACAATCTTAC	9720
TGCTCTCAAG CGGAATACCG TCATCAGAGA AACCAACCGC ACCAGCTTCT AAGAGTGCCT	9780
TAAAGTCAGT CAAGTTTTA CCATTAAGT TTTAGTAAT GGTCGCAACT GTCTTGACAT	9840
TAATCTTCTC TTTGGCAGCT GACTGGAGAA CTGCTTGCAA AGTCTCCACG TCTGAAATGG	9900
TTGGACTGGT ATTAGCCATC ATGACGACAG TAGTAAAACC ACCTGCAGCG GCTGCTAGGG	9960
CACCAAGTATG AATGTCTTCT TTATGTGTTT GACCAGGTTC ACGGAAATGA ACATGAATAT	10020
CGACCAAGCC AGGAGCAACC ACAAGACCAAG TAGCATCAAT CGTTTCTGCT CCTTCTTCCG	10080
TGATCTCAGA CGCAATTTC ATAATTTTCC CATCTTGAAC TAAGACATCA CAAACTTGAT	10140
CCAAACCAGA CTTGGGATCC ATTACACGAC CATTTCGAT TAGTAGCATC TGCTTCTCC	10200

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TTTATTCTATA GAAATCAACT TGGGTATCCA ACAATTATC CCCATCATAA ACAAACTTGG	10260
CTGAAAAGAA GGGTTTATCC TCTAAAAGCC ACTCAACAAA GGTGTGGTCA CCTTCCCAAG	10320
TCGGCTTGCT CAAAACCTCA TCATAGGAA CCCATTCTAG CGTCCCCTCA TTGCAGTCAA	10380
TCAAGTCGCC CTCAAACTCC GTCAACCTTAA AAACATAGGT GTACCAGTCT AAATCTGGT	10440
TAAATTCAAG AAAAGTGTG ACACCTTTA GAACTGGCTT GGCTTTGAGC CCTGTTCTT	10500
CAAGGATTTC ACGCGCCGCG CATTCCCTGGG GCGTCTCTCC TCTCTCTAGC TTACCACCCA	10560
CACCAATCCA TTTCCCTTCA TGGACATCAT TGGGTTTCTT ATTACGATGG AGCATGAGCA	10620
GTTCTTCCC ATTATCAATG TAGCAAATCG TCGCTAACTG AGGCATATT TCTCCTTATC	10680
TAAGCCAATC GATTGGCTCT TGTCCGTCT CTTTAAGAA TGCATTGGCC TTGGAAAAGG	10740
GCTTGGAACCC CCAAAATCCT CTATAAACCG ACAAAAGGACT TGGATGGGCT GATTCGATAA	10800
TCAAGTGATG AGGATTGGTA ACTAATGCCT TCTTCTTACG TGCTAAAGCT CCCCAGAGTA	10860
CAAAAACGAC TGGTCTATCT AGATGATTGA CCACCTGAAT CACAGCATCA GTAAAAGGCT	10920
CCCAGATTG ACCAGCATGA CCATTGGCCT GTCCAGCAGG AACAGTCAAA CAAGCATTAA	10980
GAAGCAAGAC TCCTTGCTCA GCCCAAGCTG TCAAATCATG AGATTCTTA ACTCCGATAT	11040
CATCTGACAA TTCTTTCAAG ATATTTGCA AGGATGGTGG AGCTGGGATA GAGTCAGGTA	11100
CAGAAAAACT CAAGCCCTGC GCTTGACCTG GTCCGTGATA GGGGTCTTGC CCTAGAATT	11160
CCACCTTAAC TTCTTCAAGC AGTGGTGTCA AGAGAGCCTG AAAAACCTT TCCTGGGTG	11220
GATAAATAAT CCCCTGAGAA TAGACCTGCT CCATAAACTG ATTGATTTTC CCGAAATAAC	11280
CCTCAGGTAA TTGCGCCTTA ATCAAAGCAT GCCAAGACGA GTGTTCCATA GCCGACTCGG	11340

(2) INFORMATION FOR SEQ ID NO: 148:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12127 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 148:

AAAAAAATAGA CTTGTTAGAC TATAATGTA GTAAGCCTAC ACAAGAAAAA TACATAGAGA	60
TAAAGGTGAT TATTATGAAA TTCAAAAAAA TGCTTACTCT TGCAAGCCATT GGCTTATCAG	120
GATTGGGCT TGTTGCCTGT GGCAATCAGT CAGCTGCTTC CAAACAGTCA GCTTCAGGAA	180
CGATTGAGGT GATTCACGA GAAAATGGCT CTGGGACACG GGGTGCCTTC ACAGAAATCA	240

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CAGGGATTCT CAAAAAAGAC GGTGATAAAA AAATTGACAA CACTGCCAA ACAGCTGTGA	300
TTCAAAATAG TACAGAACGGT GTTCTCTCAG CAGTTCAAGG GAATGCTAAT GCTATCGGCT	360
ACATCTCCTT GGGATCTTTA ACGAAATCTG TCAAGGCTTT AGAGATTGAT GGTGTCAAGG	420
CTAGTCGAGA CACAGTTTTA GATGGTGAAT ACCCTCTCA ACGTCCCTTC AACATTGTTT	480
GGTCTTCTAA TCTTTCCAAG CTAGGTCAAG ATTTTATCAG CTTTATCCAC TCCAAACAAG	540
GTCAACAAGT GGTACACAGAT AATAAAATTAA TTGAAGCTAA AACCGAAACC ACGBAAATATA	600
CAAGCCAACA CTTATCAGGC AAGTTGTCTG TTGTAGGTT CACTTCAGTA TCTTCTTTAA	660
TGGAAAAATT AGCAGAACGCT TATAAAAAG AAAATCCAGA AGTTACGATT GATATTACCT	720
CTAATGGGTC TTCAGCAGGT ATTACCGCTG TTAAGGAGAA AACCGCTGAT ATTGGTATGG	780
TTTCTAGGGA ATTAACCTCCT GAAGAACGGTA AGAGTCTCAC CCATGATGCT ATTGCTTTAG	840
ACGGTATTGC TGTTGTGGTC ATAATGACA ATAAGGCAAG CCAAGTCAGT ATGGCTGAAC	900
TTGCAGACGT TTTTAGTGGC AAATTAACCA CCTGGGACAA GATTAAATAA AATGTTGCT	960
CCATAAAATCT CTAAAGAGAT GCAGACGTTT CATCGTACAA TAAGATAAAG AAGGCAAGTA	1020
GGGAGGTGTC GTATCTCCCT TACTTTCTTC ACTAGAAAGG ACAAGATGTG ACAAAACAAG	1080
CCTTCAAAGA AGCAGTTTTT AGGGCAATT TTTCATGAG TGCAACAGTA GCTGTTGTAG	1140
CTATTTGCT AATCTGTTTC TTTATTTTTA GTAATGGCTT ACCTTCATA GCTAACTACG	1200
GCTTGGCCCG TTTTTTATTAA GGCAAGTGATT CGTCGCCAAC GAACATTCCG GCAAGCTATG	1260
GTATTTTACCA AATGATCGTT GGTTCCATTAT TAATTACCTT AGGAGCGATT GTGATTGGGG	1320
TGCAACACAGG CATCTTGACA TCGGTGTTA TGTTTATTAA TTGTCAAAG CCCGCTATG	1380
GCTTCTTAAA ATCAGCTATC AACTGATGG CAGCCATTCC ATCTATTGTT TATGGTTTTT	1440
TCGGCCTACA ATTATTGGTG CCTTGGATTA GAAGCTTTT AGGAAATGGC ATGAGTGTCC	1500
TAACCGCTTC GTTACTATTAA GGAATAATGA TTTTGCCAAAC CATTATCAGT TTGTCAGAAT	1560
CTGCTATCCG AACAGTTCCC AAAACGTATT ATTCTGGTAG CTTGGCTCTA GGAGCTAGTC	1620
ATGAACGGAG TATTTTTAGT GTCATCTTGC CAGCTGCGAG ATCTGGTATT TTATCAGCAG	1680
TTATTTAGG AATCGGTCGC GCAGTAGGTG AAACCATGGC AGTTATTTG GTGGCAGGCA	1740
ACCAGCCGAT TATTCCAAGT GGACTCTTTT CAGGAACCAG AACCTTAACA ACCAATATTG	1800
TTCTGGAAAT GGCTTACGCA TCAGGTCAGC ATAGGGAAGC CCTTATTGCA ACCTCAGCAG	1860
TTCTCTTTT CCTTATTCTC TTGATTAATG CCTACTTTGC CTACTTGAAA GGAAAATCAT	1920
CTTATGAGTA AATACCTGCT AAAACTCTC GTTTATTGTT TTTCAGCTTT AACCTTGGC	1980
TCTCTCTTT TAATCATTGG TTTTATCCTC ATCAAAGGCT TACCTCATCT AAGTCTATCC	2040

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CTCTTTCTT GGACTTATAC TTCTGAGAAC ATTCCCTTA TGCCAGCGAT TATTCCACC	2100
GTTATTCTGG TCTTGGTGC TCTTCTTTA GCCTTGCCCA TAGGGATTTC TGCTGGTTT	2160
TATCTGTGG AATATACAAA AAAAGATTCC CTTGTGTTA AAATCATGCG ATTGGCCTCA	2220
GATACCTTAT CTGGGATTCC TTCCATTGTT TTTGGTCTGT TTGGCATGCT CTTCTTGTA	2280
GTCTTCTTAG GTTTCAATA CTCTCTGTTA TCAGGAATCT TAACCTCAGT TATCATGGT	2340
TTGCCAGTCA TTATTCGCTC AACAGAAGAA GCCCTTTAT CTGTTAGTGA TAGCATGCGT	2400
CAAGCAAGTT ATGGACTTGG GGCAGGTAAG TTACGGACTG TTTTAAAGAT TGTTCTACCA	2460
GTTGCCATGC CAGGTATTT AGCTGGAGTG ATACTAGCTA TTGGCCGTAT CGTTGGTGAA	2520
ACAGCTGCC TCATGTATAC ATTAGGTACC TCTACCAATA CGCCAAGTAG TCTCATGTCT	2580
TCAGGCCGTT CTCTAGCCCT ACATATGTAT ATGCTGCAA GTGAGGGGCT ACATGTCAT	2640
GAAGCCTATG CTACCGCCGT GATTTTGATT ATTACTGTTT TAATGATAAA TACTCTATCA	2700
AGCTTATTAT CTCGAAAATC TGTGAAAGGA GCTTCCTAGT ATGGAAACAT TTTCAGTCAG	2760
ACACCTAGAC TTATTTACG GGGATTTCA AGCCTAAAAA AATATTCGA TTCAATTACC	2820
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AACCTTAAC CGGATGAACG ATTTGGTCC TTCTTGCCAT ATTGAAGGCC AAGCCTCTT	2940
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GATGGTTTT CAACAGCCTA ATCCCTTGCA CATGTCTATC TATGATAACG TGGCTTATGG	3060
CCCAAGGACA CATGGTATTC GAGACAAAAA ACAATTAGAT GCCTTAGTGG AGAAATCTT	3120
AAAAGGGGCA GCCATTTGGG AAGAAGTCAA AGATGATCTT AAAAAGAGTG CCATGTCCTT	3180
ATCTGGCGGT CAGCAGCAAC GCCTTGCAAT TGCGCGAGCT TTAGCAGTAG AACCTGATAT	3240
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CCTCATTCAAG CAACTAAAAA AGGATTATAC GATTATCATT GTTACCCATA ACATGCAACA	3360
AGCTTCACGT ATTCAGATA AAACTGCTTT TTTCTTAACA GGAGAAATTG GCGAATTGG	3420
AGATAACCGTT GACGTGTTA CCAATCCAAA AGATCAGCGC ACAGAAGACT ATATTCAGG	3480
ACGGTTCGGA TAAGGAAGGA AAAACCTATG AGAAATCAAT TTGACTTAGA ATTGCATGAA	3540
TTAGAACAAAT CCTTTTAGG ACTAGGGCAA CTGTCCTTG AAACAGCTTC AAAAGCCTTA	3600
CTGGCCTTAG CCTCCAAAGA CAAGGAGATG GCAGAGCTAA TTATCAATAA GGATCATGCT	3660
ATCAACCAAG GTCAAAGCGC TATCGAATTG ACCTGTGCC GTTGTGTC CTTGCAGCAG	3720
CCACAAGTGT CTGACCTTCG ATTTGTGATT AGCATCATGT CTTCTTGTC AGACCTTGAA	3780

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CGTATGGGAG ACCATATGGC AGGCATTGCC AAAGCTGTT TGCAACTAAA AGAAAATCAA	3840
CTAGCCCCCTG ACGAAGAACAA GTTACACCAA ATGGGTAAAT TATCCCTCAG CATGCTAGCC	3900
GATTTATTGG TTGCCTTCC TTTGCACCAA GCCTCAAAAG CTATTAGTAT TGCTCAAAAA	3960
GATGAACAGA TTGACCAATA TTATTATGCC TTATCAAAGG AAATCATGG ACTTATGAAA	4020
GACCAAGAAA CCTCAATTCC CAATGGAAC CAATACCTT ATATCATAGG GCATCTGGAA	4080
CGCTCGCTGA TTACATTGCT AACATTTGTG AACGCCTAGT CTACCTAGAA ACAGGAGAAC	4140
TAGTGGATTG GAATTAATTC AACTAATCCT TAAAAGAGAA GAGTACGATT AAGTACTCTT	4200
TTTTATGGTT GTAAAAAAAGT TCATTTGACC AATTAAAGCA GTGTAGATAG TGAGGAGTTG	4260
TTTCAATTCT ATCGTGAACG AGGGAAATGCT GAAAACCTTA TCAAAGAAAG GAAAGCAGGA	4320
TTCTTTGGGG ATAAGACAGA TAGTCGACC ATGATTAAGA ATGAAGTACG TATGATGATG	4380
GGCTGTCTGG CTTATAATCT CTACCTCTT TTAAAGCAGC TAGCTGGTGA TGAAGTAAAG	4440
TCCTTGACTA TCAAGCGTTT TCGACGTCTC TTCCCTCATA TTGCCGGAAA ATATGCTCT	4500
ACTGCTAGAC GACATATTCT CAAATTCTCA AGTCTATACG CCTATTCAAACAGTTCAA	4560
GCCTTATTG ATACAATCTG CCAGATAAAAT CTGATACTCC CTGTTCCATA TAGAGCTAGA	4620
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CCAAGGGAGG AGTCTGCCCT TTTTTAGGAA AAAATCAAGA CAAATCTCCT CAATTATGTC	4740
TCGAACATCA GAAATTAAGC AAAATCACCA GAAGGACAGT ATTTCAACTA GCTTTCTGG	4800
TAATTTTTGA ACTGTGTAGT TCGTTAGTGC CAGATATGAA TAATTGGGTA TGATAAAATCT	4860
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TAGGTATTCC TTATCCAACCT CTATATAACT TGGCATCAAC TTGTAATCTT CAACCCCCAA	5040
ACGTTCAGCA ATATATTTTA ACTTTGTTAG TATTGGTCTG GATTCTCCAT TTTCAAATTCT	5100
AATTAATTGA CGGATACTTA ATTCAGACTC ATCACCACAA AATTCTGAAC GACTGATT	5160
TTTAGCCAAA CGTAATCTTT TAATTTTTTC GCCAAACTCT CGCAACCTAC AAGAACTTCC	5220
TGAGTTGTTT ACCTCTATTA TAAGCATATA CTGAATCAA CTATCTATCA GATTCTTCT	5280
CACTTTAACT AAAGACTAAG AGTTTATCCC TTCGTCTCGG TTTTTGTGTA TTTTCCACC	5340
ATACCCAGT AATGCAAGTG CAAAATCCCC TAGAATATGA TAGAATAAGA GAAAGAACTC	5400
TATCAAGGAG GAAATCATGG AAAAACAAAC CGTCGCCGTC TTGGGGCCTG GTTCTTGGGG	5460
AACCGCCCTT TCACAAGTCT TAAATGACAA TGGACACGAG GTACGTATTT GGGGAAATCT	5520
TCCCGAGCAA ATCAATGAAA TTAATACACA CCATACTAAT AAGCACTACT TTAAAGATGT	5580

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CGTTCTAGAC GAAAATATCA TTGCCTACAC CGACTTAGCA GAAACATTGA AAGATGTGGA	5640
TGCGATTTG TTTGTTGTC CAACAAAAGT GACACGACTT GTTGCCCAGC AAGTTGCACA	5700
AACCTTGGAC CATAAGTTA TCATCATGCA CGCATCAAAG GGATTAGAAC CTGATAGCCA	5760
TAAACGATTA TCAACCATTG TTGAAGAAGA AATTCCCTGAA CATCTCCGTA GTGATATCGT	5820
CGTTGTTCA GGGCCTAGTC ATGCAGAAGA GACCATTGTG CGTGACCTAA CTTTAATAAC	5880
TGCTGCTTCT AAAGATTAC AAACAGCTCA ATACGTTCAAG AAGCTATTAA GTAATCACTA	5940
CTTCCGACTT TATACCAATA CGGATGTTAT CGGGGTTGAA ACTGCTGGTG CTCTTAAAAA	6000
TATTATTGCT GTCGGTGCTG GAGCTTACA TGGCTTGGA TTTGGTGATA ATGCTAAGGC	6060
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TCCATTGACC TATAGCGGCT TATCTGGTGT GGGAGATTTG ATCGTAACGG GAACTTCCAT	6180
CCACTCTCGT AACTGGAGAG CTGGAGATGC TCTCGGACGA GGAGAATCCC TAGCTGATAT	6240
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AGCCAAGAA CTTGGAGTCT ATATGCCAT TACACAGGCT ATTTACCAAG TTATTATCA	6360
CGGAACCAAT ATCAAAGATG CCATTTATGA CATCATGAAC AATGAATTAA AAGCAGAAAA	6420
TGAGTGGTCT TAACCCCTCA TAGAAAGGAT TTTTATGACA TCAAAAGTTA GAAAGGCAGT	6480
CATCCCTGCT GCTGGACTAG GAACTCGATT TTTACCAGCA ACCAAGGCC TTGCCAAAGA	6540
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TGATTCAAC TTCGAATTGG AATATAACCT CAAAGAAAAA GGGAAAACAG ATCTTTGAA	6720
GCTAGTTGAT AAAACAAC TG ACATGCGTCT GCATTTTATC CGCCAAACTC ATCCACGCGG	6780
TCTCGGAGAT GCTGTTTGC AAGCCAAGGC TTTCGTCGGA AATGAACCTT TTGTCGTTAT	6840
GCTTGGTGAT GACTTGATGG ATATCACAGA CGAAAAGGCT GTTCCACTTA CCAAACAACT	6900
CATGGATGAC TACGAGCGTA CCCACGCGTC TACTATCGCT GTCATGCCAG TCCCTCATGA	6960
CGAAGTATCT GCTTACGGGG TTATGCTCC GCAAGGCGAA GGAAAAGATG GTCTTACAG	7020
TGTTGAAACC TTTGTTGAAA AACCAAGCTCC AGAGGACGCT CCTAGCGACC TTGCTATTAT	7080
CGGACGCTAC CTCCTCACGC CTGAAATTGG TGAGATTCTC GAAAAGCAAG CTCCAGGTGC	7140
AGGAAATGAA ATTCAAGGGG CTCGTTACGA TGTGGAGAC AAGTTGGCT TCATGAAAC	7200
TGCTCGTGAG TTCAAAGGGG CTCGTTACGA TGTGGAGAC AAGTTGGCT TCATGAAAC	7260
ATCCATCGAC TACGCCCTCA AACACCCACA AGTCAAAGAT GATTGAAGA ATTACCTCAT	7320

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CCAACCTGGAA AAAGAATTGA CTGAGAAGGA ATAACAAAAT CATTATATA AAGATTAGCC	7380
ACACATAAAAT TAAGTAAATT CTCTACTTGA ATCTACCTAT TTAATAAAAA CTAATGAAAA	7440
CGCTATACTT GTATTTGTTT TTTCATTAAA ATAAGAGTAG AATAAATTAG TATAGTAAA	7500
CAAAAAGCA CCGAACCGGT GCGCACTTT TCAAGTTGTG TACGGACAAA GCCTTATTT	7560
AACTTGCTA TGTTGTTCT AATGGTTCCA AAATAATAAA TAATTTAAA TTTGACTTAA	7620
CTGTTGGAGT AGTCATGGTT AAATAAACC AACCGAGCCG AACATAAGTT GTTTAATT	7680
GTGGAAGCTA TTAATAAAAA TATAATAAGG GAGAAAGATA GGTGTAATT TAATTTAAA	7740
GTAATTGCAG ACACATATCAA AGAAAAAGAT TATGGAGAAC AAATTTGTAG AATTATCGA	7800
AAACAATAAA AAAGTAATCA TTTCATCAGT TCCAGTTGGT GTTGTATTGG TATTAGGGTT	7860
TGGATGGTAT TCATATAACC AACACAACGC AGAACACAA GCAAAATTG TACAATTAGA	7920
AAAAGATAGC AAATCAGACA AAGAACAAAGT TGATAAACTA TTGATCAT TTGATGCATC	7980
TTCAGATGAA TCTATTTCTA ATTAAAAGA ACTATCTGAA ACTTCACTTA AAACCGATGC	8040
AGGTAAAGAC TATCTTAATA ACAAAAGTCAA AGAACATCT AAAGCAATTG TAGATTTCA	8100
TTTGCAAAAA GGTTTGGCTT ATGATGTTAA AGATTCAAGAT GACAAATTAA AAGATAAAAGC	8160
AACTCTTGAA ACAAAATGTA AAGAAATTAC AAAACAAATT GATTTTATCA AAAAGATTGA	8220
TGAAACTTTT AAACAAGAGA ATTTGGAAGA AACTCTTAAA TCTCTAAATG ATCTTGTGA	8280
TAAATATCAA AAACAAATCG AACTTTGAA GAAAGAAGAA GAAAAAGCTG CTGAAAAGC	8340
TGCTGAAAAA GCAAAGGAAT CTTCTAGTCA AAGTAATTCT TCTGGTAGTG CTTCTAATGA	8400
GTCTTATAAT GGATCTTCCA ATTCAAATGT AGATTATAGT TCATCTGAAC AACTAATGG	8460
ATATTCAAAT AATTATGGCG GTCAAGATTA TTCTGGTTCA GGAGATAGTT CAACAAATGG	8520
TGGATCATCA GAACAATATT CATCTAGCAA TTCAACAGC GGAGCAAATA ATGTCTACAG	8580
ATATAAAGGC ACTGGTGCTG ACGGCTATCA AAGATACTAC TACAAAGATC ATAATAATGG	8640
AGATGTGTAT GATGACGATG GAAATTACCT TGGGAACCTT GGTGGCGGCA TTGCAGAAC	8700
TAGTCAACGCC TAATAACTAT TTTAGAGCTG TGGTGTTCG AATGGTTCCA AAAACACATTA	8760
AAAGCTACTC ATTTTTAAG TAGCTTTTT CTTATTCAAG TTTACATATT ATACTCAATG	8820
AAAATCAAAT TCAAACCAAG TCAGCATCGC CTTACCGTAG GTATGGTTAC TGACTTCGTC	8880
AGTTTCATCT ACAACCTCAA AACCATGTT TGAGCTGACT TCGTCAGTTC TATCTACAAC	8940
CTCAAAGCAG TGCTTGAGC AACCTGCGGC TAGCTTCCTA GTTTGCTCTT TGATTTCAT	9000
TGAGTATTAG TCGTCACAAAT CCCATTCCCT TGTAGAAAAG CAAAATGGCG AGTCCTACGA	9060
ACAAGACTAC CGCTCCTAAT CTCTGGCTGG TGTTATACAT CCGTTTTCT CCTCTAACTG	9120

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GAAAGATAAC	TGCTAGAACAT	GCGCCACCAA	CTGCACCACC	GATATGGCCT	GCTAGGCTGA	9180
TTCCTGGAAT	CAGAACACTT	CCAATAATGT	TAACCACAAA	AAGTGTCA	AG TAGGATTGCC	9240
CTAGCTGTTG	GATATAAGGA	TTGCGAGTTG	CATAGCGAAG	AACAATAATC	GGCGCAAATA	9300
GCCCATAAAAG	AGAGGTAGAG	GCGCCTGCTG	CTAAGGATT	AGGACTAAAT	ACAAAAACAA	9360
AGAGATTGCC	CATCATTCCT	GATAAAAGAT	AGAGAAAGAA	AAACTGCTTA	GAACCGAAAA	9420
TCTCCTCTAC	CTGCCTTCCA	AGATAATAAA	GTGAAAGCAT	ATTAACAATG	AAATGTTCCC	9480
ACCCAATATG	AACAAAAATG	GCAGACAAGA	GACGCCAAC	CTGCTCGGG	AAGAGGCGAA	9540
TAGCTGGCCC	ATACATGGCT	CCAAATCGAA	ATAATGTATC	TGCCCTGTCA	AAGTTTCCGC	9600
CTGCAGTGAC	CAACATTAGT	AAAAATACCA	AGGCCGTAC	TAAGAGGAAG	AAACTCGTCA	9660
CAGGGTAACG	TCTATCAAAG	ATTTCCCTCA	TCAATTATA	CCTCCTGAAC	AGGAATATCA	9720
TGGTTTCAG	GTATAAAGTC	CTGAATTG	CAAGGATATA	TCGTA	CTCAA AGTACCGACCA	9780
GAAAAATGTT	CCAGATAGCG	GTCATAATAG	CCTCCACC	GT ATCC	CTATCCG ATATCCTT	9840
GTCGTAAAG	CCAGACCAGG	AACATGAATC	AAATCAATCT	GAGATGCATC	CACCACTTCC	9900
AAATCTCCCT	GTAGCTCCAG	TAAGGCAAAG	AAAGTTTTA	CCAACTGTTG	CGGATCATAG	9960
ACCACAAAGT	CCATGCGCCC	CTTGGGATAA	GT	TTGGT	TTAAACCTT CTTGCCGTCC	10020
TTCAGCGCCT	GCTCAATCAG	TTCTGCGTT	TGAAACTCAT	GAGAAAAGA	GAGGTAGGTT	10080
GCGATGACCT	TGGCTCTTG	ATAAAAGGGG	TGTTGTAAAA	GCCGCTCGGT	TAAAGCTTGG	10140
TCTATAGCCT	GT	TTTGCTC	TTGAGATATA	GCCTTCATT	CATGCAAGAC TTGCTTGCGT	10200
AATTCCGATT	TCATAGACAA	GCCCTCTATT	CTGCTGCCTT	CTTTTTCA	GG AAAACTAGACA	10260
CCGCAGCCAC	CCCAATAGCT	AAGACTTCTT	CCTTAGGACT	CATT	TGAGGG TGATGAAGAG	10320
CGTAGGGACT	ATCGATACCT	AGCCAAAACA	TCACGCCATC	AACCTTGAA	AGGAGATAAC	10380
CAAAGTCCTC	GCCTGTCATA	GCAGGTTCGA	TATCAATCAA	CTCGATTCCG	TCTTTTCGT	10440
CAAAGAAAGTC	CATCAGTTCA	CGCGCCAAGG	CTGGATTGTT	CTCAACAGGT	AGGTATCCAC	10500
CTTGTGAG	TTCCACTTCG	ACTTCCATAT	CAAAGGCAGC	TGCAACCCCT	TCTGCAACTG	10560
TTTTTACCCCT	CTTTTGCA	AAGAGACTCA	TGTCCTGTGT	CAAGGCACGA	ATAGTTCCAT	10620
GTAAAAAAAGC	TGTGTCTGTG	ATGACATTGT	TGGTGGTTCC	AGCTTGAAA	ACGCCGAAGG	10680
TCACCACTGC	TCCCTCGATT	GGGTTGACAT	TGCGGCTAAC	AACTGACTGC	ACTTGGGTCA	10740
CAAAGTAACT	AGCCGCCACC	AAGGCACAT	TGGCTTCATG	AGGAAAAGCT	GC GTGGCCAC	10800
CTTTGCCTTT	GAAACGGATC	TTCACCTCGC	AAAGTCCTGC	AAAGAGTGT	TA TGAGTATTAG	10860

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TCGCAATCTG	GCCGACTTTC	AAATCTGGAC	GAACATGGAG	ACCATAGAAT	TGATCTGGCA	10920
ACCAATCTCC	AAAAGCACCG	TCCTCATACA	TGAGCATACC	ACCAGCTTCA	TTTTCTTCAG	10980
CAGGCTGAAA	TAGAAAGAGC	AGATTATTCT	TGGGTTGCTC	CTCAAGGGCG	CGCTCAAGAC	11040
AGCCTAAGGC	AATGGTCATA	TGAAAATCAT	GGACACAGGC	ATGCATGCGA	CCTTGGTGTT	11100
GAGAAGCAAA	AGGTAGACCT	GTGGTTCGA	CGATAGGCAG	GCCATCAATA	TCTGTCCGCC	11160
AACCAATGGT	TCGCTCCGGC	TGACTTCCT	GCAGGGTAGAC	CAAATCCCT	GTCCGCCAAG	11220
TACGAATTTG	AACAAAATCC	TTGCCCGTAG	TCAATTTCTC	AATCACATCC	AGCAAATAAG	11280
CCTGAGTCTT	GAACCTCTCC	AAGCCAATCT	CTGGAATCTG	GTGTAAATCT	CGTCTAGTCT	11340
GAATCAAATC	TAACATCTAT	CTGTCCTCCG	ATATAGCAGA	AAGAGGCTGG	AAAAAGGGTT	11400
CCGCCTCTTT	TTTACTTTA	CAATTACAAG	GTACGAAGCG	CATCCTCTAG	CGCTGTTTT	11460
TGTTGAGTTT	GGGCATCAAT	TTCTTGATA	ATACGAGCTG	GAACACCTGC	TACTACCACG	11520
TTTTCTGGGA	CATCTTGGGT	AAACAATAGCT	CCTGCTGCGA	CAACTGAACC	ACTACCGATT	11580
TGGACTCCTT	CGATAACCAC	TGCATTAGCA	CCGATAAAGAA	CATTGTCTCC	GACACGGACT	11640
GGTCAGCAC	TAGCTGGCTC	AATCACACCT	GCCAAAACGT	CACCTGCACC	AACGTGGCTA	11700
TTTTTCCCAA	CGATGGCACG	GCCACCAAGG	ATGGCACCCA	TGTCAATCAT	GGTCCAGCA	11760
CCGATTTCA	CACCGATATT	GATAACAGAT	CCCATCATGA	TAACAGCATT	GTCACCAATT	11820
TCCACCTGGT	CACGGATAAT	CGCACCTGGC	TCGATACGAG	CGTTGATAGC	ACGCTTATCT	11880
AGCAAAGGAA	CTGCAGAATT	ACGAGCATCT	TGCTCGACAA	CATAATCTTG	ATTTCTTAC	11940
AAACCTTCAA	GAAGCGGAGC	CACATCCTTC	CAGTCTCCGA	ATAGGACATT	TCCTAGTTG	12000
ACAACAGAGC	TAGGCACAGC	AGTTGCGAGT	TGCCCCTCAA	AGGTTACTTT	GACACTGGTT	12060
TTCTTTTCAG	CATTGGCGAT	AAATTGGATA	ATTTCATTGAG	CGTTCATTTC	TGTAGCAGTC	12120
ATAGGTG						12127

(2) INFORMATION FOR SEQ ID NO: 149:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 12566 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 149:

CCATCCTTCT	GTTGATGTGA	CAGGAATGAT	GATAATCAA	CCAGTAGCTA	GTCGCGAAGA	60
GGTGACAGAG	GCTTGAGTC	ACTTGGCGGT	AGAGCACAAT	AGTCTCATTG	CTCGTCGAAT	120